(19) World Intellectual Property Organization

International Bureau





(43) International Publication Date 25 March 2004 (25.03.2004)

PCT

(10) International Publication Number WO 2004/024162 A1

[DE/DE]; Wingertstrasse 24, 69207 Sandhausen (DE).

HERMANN, Kristina [DE/DE]; Viernheimer Strasse 32,

(74) Agents: KRAUSS, Jan, B. et al.; Boehmert & Boehmert, Pettenkoferstrasse 20-22, 80336 Munich (DE).

(81) Designated States (national): AE, AG, AL, AM, AT, AU,

(51) International Patent Classification⁷: A61K 31/517, C07D 239/95, 401/12, 403/04, A61P 3/06

(21) International Application Number:

PCT/EP2003/007067

(22) International Filing Date: 2 July 2003 (02.07.2003)

(25) Filing Language: English

(26) Publication Language: English

(30) Priority Data:

02020255.2 10 September 2002 (10.09.2002) EP

AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, OM, PH, PL, PT, RO, RU, SC, SD, SE, SG, SK, SL, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, YU, ZA, ZM, ZW.

(71) Applicant (for all designated States except US): LION BIOSCIENCE AG [DE/DE]; Waldhofer Strasse 98, 69123 Heidelberg (DE).

(72) Inventors; and

(75) Inventors/Applicants (for US only): DEUSCHLE, Ulrich [DE/DE]; Wundtstrasse 9/20, 69123 Heidelberg (DE). LOEBBERT, Ralph [DE/DE]; Hildastrasse 21, 69115 Heidelberg (DE). BLUME, Beatrix [DE/DE]; Heinrich-Heine-Strasse 20, 69221 Dossenheim (DE). KOEGL, Manfred [DE/DE]; Hauptstrasse 131/4, 69214 Eppelheim (DE). KREMOSER, Claus [DE/DE]; Mozartstrasse 29, 69121 Heidelberg (DE). KOBER, Ingo [DE/DE]; Am Grossen Wald 30, 69251 Gaiberg (DE). BAUER, Ulrike (84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZM, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HU, IE, IT, LU, MC, NL, PT, RO, SE, SI, SK, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

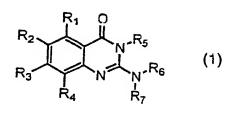
Published:

with international search report

68623 Lampertheim (DE).

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

54) Title: 2-AMINO-4-QUINAZOLINONES AS LXR NUCLEAR RECEPTOR BINDING COMPOUNDS



(57) Abstract: The present invention relates to 2-amino-4-oxo-quinazolines according to the general formula (1), which bind to the LXR receptors and act as agonists and antagonists of the LXR receptors. The invention further relates to the treatment of diseases and/or conditions through binding of said nuclear receptor by said compounds and the production of medicaments using said compounds. In particular the compounds are useful in the treatment of hypercholesterolemia, obesity or other diseases associated with elevated lipoprotein (LDL) levels.

WO 2004/024162 PCT/EP2003/007067

2-AMINO-4-QUINAZOLINONES AS LXR NUCLEAR RECEPTOR BINDING COMPOUNDS

The present invention relates to compounds according to the general formula (1), which bind to the LXR receptors and act as agonists and antagonists of the LXR receptors. The invention further relates to the treatment of diseases and/or conditions through binding of said nuclear receptor by said compounds and the production of medicaments using said compounds.

BACKGROUND OF THE INVENTION

Multicellular organisms are dependent on advanced mechanisms of information transfer between cells and body compartments. The information that is transmitted can be highly complex and can result in the alteration of genetic programs involved in cellular differentiation, proliferation, or reproduction. The signals, or hormones, are often simple molecules, such as peptides, fatty acid, or cholesterol derivatives.

Many of these signals produce their effects by ultimately changing the transcription of specific genes. One well-studied group of proteins that mediate a cells response to a variety of signals is the family of transcription factors known as nuclear receptors, hereinafter referred to often as "NR". Members of this group include receptors for steroid hormones, vitamin D, ecdysone, cis and trans retinoic acid, thyroid hormone, bile acids, cholesterol-derivatives, fatty acids (and other peroxisomal proliferators), as well as so-called orphan receptors, proteins that are structurally similar to other members of this group, but for which no ligands are known (Escriva, H. et al., Ligand binding was acquired during evolution of nuclear receptors, PNAS, 94, 6803 – 6808, 1997). Orphan receptors may be indicative of unknown signaling pathways in the cell or may be nuclear receptors that function without ligand activation. The activation of transcription by some of these orphan receptors may occur in the absence of an exogenous ligand and/or through signal transduction pathways originating from the cell surface (Mangelsdorf, D. J. et al., The nuclear receptor superfamily: the second decade, Cell 83, 835-839, 1995).

In general, three functional domains have been defined in NRs. An amino terminal domain is believed to have some regulatory function. A DNA-binding domain hereinafter referred to as "DBD" usually comprises two zinc finger elements and recognizes a specific Hormone Responsive Element hereinafter referred to as "HRE" within the promoters of responsive genes. Specific amino acid residues in the "DBD" have been shown to confer DNA sequence binding specificity (Schena, M. & Yamamoto, K.R., Mammalian Glucocorticoid Receptor Derivatives Enhance Transcription in Yeast, Science, 241:965-967, 1988). A Ligand-binding-domain hereinafter referred to as "LBD" is at the carboxy-terminal region of known NRs. In the absence of hormone, the LBD of some but not all NRs appears to interfere with the interaction of the DBD with its HRE. Hormone binding seems to result in a conformational change in the NR and thus opens this interference (Brzozowski et al., Molecular basis of agonism and antagonism in the oestrogen receptor, Nature, 389, 753 – 758, 1997; Wagner et al., A structural role for hormone in the thyroid hormone receptor, Nature, 378, 690 – 697. 1995). A NR without the HBD constitutively activates transcription but at a low level.

Coactivators or transcriptional activators are proposed to bridge between sequence specific transcription factors and the basal transcription machinery and in addition to influence the chromatin structure of a target cell. Several proteins like SRC-1, ACTR, and Grip1 interact with NRs in a ligand enhanced manner (Heery et al., A signature motif in transcriptional coactivators mediates binding to nuclear receptors, Nature, 387, 733 – 736; Heinzel et al., A complex containing N-CoR, mSin3 and histone deacetylase mediates transcriptional repression, Nature 387, 43 – 47, 1997). Furthermore, the physical interaction with repressing receptor-interacting proteins or corepressors has been demonstrated (Xu et al., Coactivator and Corepressor complexes in nuclear receptor function, Curr Opin Genet Dev, 9 (2), 140 – 147, 1999).

Nuclear receptor modulators like steroid hormones affect the growth and function of specific cells by binding to intracellular receptors and forming nuclear receptor-ligand complexes. Nuclear receptor-hormone complexes then interact with a hormone response element (HRE) in the control region of specific genes and alter specific gene expression.

The term LXR (Liver X Receptor) includes all subtypes of this receptor. Specifically LXR includes LXRa (also known as LXRalpha, RLD-1 and NR1H3) and LXRb (also known as LXRbeta, NER, NER1, UR, OR-1, R1P15 and NH1H2) and ligands of LXR should be under-

stood to include ligands of LXRa or LXRb. LXR is a prototypical type 2 nuclear receptor which activates genes upon binding to promoter region of target genes in a prototypical heterodimeric fashion with Retinoid X Receptor (hereinafter RXR, Forman et al., Cell, 81, 687-93, 1995). The relevant physiological ligands of LXR seem to be oxidized derivatives of cholesterol, including 22-hydroxycholesterol and 24,25(S)-epoxycholesterol (Lehmann, et al., Biol. Chem. 272(6), 3137-40, 1997). The oxysterol ligands bound to LXR were found to regulate the expression of several genes that participate in cholesterol metabolism (Janowski, et al., Nature, 383, 728-31, 1996).

LXR is proposed to be a hepatic oxysterol sensor. Upon activation (e.g. binding of oxysterols) it influences the conversion of dietary cholesterol into bile acids by upregulating the transcription of key genes which are involved in bile acid synthesis such as CYP7A1. Hence, activation of LXR in the liver could result in an increased synthesis of bile acids from cholesterol which could lead to decreased levels of hepatic cholesterol. This proposed LXR function in hepatic cholesterol metabolism was experimentally confirmed using knockout mice. Mice lacking the receptor LXRa lost their ability to respond normally to an increase in dietary cholesterol and did not induce transcription of the gene encoding CYP7A1. This resulted in accumulation of large quantities of cholesterol in the livers and impaired hepatic function. (Peet, et al., Cell, 93, 693-704, 1998).

Besides its important function in liver, LXR plays an important role in the regulation of cholesterol homeostasis in macrophages and intestinal mucosa cells where it upregulates cholesterol transporters from the ABC (=ATP binding cassette) family of membrane proteins (Repa, et al., J Biol Chem. 2002 May 24;277(21):18793-800). These transporters are believed to be crucially involved in the uptake of cholesterol from the diet since mutations in their genes leads to diseases such as sitosterolemia (Berge, et al., Science (2000);290(5497):1771-5.).

Other members of the ABC transporter family seem to be responsible for the efflux of cholesterol from loaded macrophages, a process which is thought to prevent the generation of atherosclerotic lesions. Stimulation of LXR by synthetic ligands might result in an increased cholesterol efflux from macrophages and a decreased deposition of atherosclerotic plaques (Venkateswaran, et al., PNAS (2000) 24;97(22):12097-102; Sparrow, et al., J Biol Chem (2002) 277(12):10021-7; Joseph, et al., PNAS (2002);99(11):7604-9).

However, in animal studies it was observed that activation of LXR in the liver by full agonists does not only increase bile acid synthesis but also stimulates the de novo synthesis of fatty acids and triglygerids through the upregulation of key enzymes such as Fatty Acid Synthase (FAS) or Stearyl-CoA Desaturase (SCD-1). (Schultz, et al., Genes Dev (2000) 14(22):2831-8.

Therefore, an ideal synthetic LXR binding compound should have properties that retain the agonistic activity on hepatic bile acid formation and ABC-transporter -mediated decrease in cholesterol uptake from the diet and increased cholesterol efflux from macrophages. In parallel such a compound should lack the hyperlipidemic potential which is exerted through increased fatty acid and triclyceride synthesis.

To date few compounds have been described which bind the LXR receptor and thus show utility for treating diseases or conditions which are due to or influenced by said nuclear receptor (Collins, et al., J Med Chem. (2002) 45(10):1963-6; Schultz, et al., Genes Dev (2000) 14(22):2831-8; Sparrow, et al., J Biol Chem (2002) 277(12):10021-7).

It was thus an object of the present invention to provide for compounds which by means of binding the LXR receptor act as agonist, antagonist or mixed agonist / antagonist of said receptor and thus show utility for treating diseases or conditions which are due to or influenced by said nuclear receptor.

It was further an object of the invention to provide for compounds that may be used for the manufacture of a medicament for the treatment of cholesterol associated conditions or diseases. In a preferred embodiment of the invention it was an object of the invention to provide for compounds that lower serum cholesterol and/or increase High Density lipoproteins (HDL) and/or decrease Low Density Lipoproteins (LDL). It was also an object of the invention to provide for compounds that may be used for the treatment of lipid disorders including hypercholesterolemia, atherosclerosis, Alzheimer's disease, skin disorders, obesity and diabetes.

SUMMARY OF THE INVENTION

The present invention provides, *inter alia*, novel LXR nuclear receptor protein binding compounds according to the general formula (1) shown below. Said compounds are also binders of mammalian homologues of said receptor. Further the object of the invention was solved by

providing for amongst the LXR nuclear receptor protein binding compounds according to the general formula (1) such compounds which act as agonists, antagonists or mixed agonists / antagonists of the human LXR receptor or a mammalian homologue thereof.

The invention provides for LXR agonists that may be used for the manufacture of a medicament for the treatment of cholesterol associated conditions or diseases. In a preferred embodiment of the invention it was an object of the invention to provide for compounds that lower serum cholesterol and/or increase High Density lipoproteins (HDL) and/or decrease Low Density Lipoproteins (LDL). It was also an object of the invention to provide for compounds that may be used for the treatment of lipid disorders including hypercholesterolemia, atherosclerosis, Alzheimer's disease, skin disorders, obesity and diabetes.

The foregoing merely summarizes certain aspects of the present invention and is not intended, nor should it be construed, to limit the invention in any manner. All patents and other publications recited herein are hereby incorporated by reference in their entirety.

DETAILED DESCRIPTION OF THE PREFERRED EMBODIMENTS

The invention provides for a compound according to the following formula (1), or pharmaceutical acceptable salts or solvates thereof, hereinafter also referred to as the "compounds according to the invention" including particular and preferred embodiments thereof, wherein

$$R_2$$
 R_3
 R_4
 R_5
 R_6
 R_7 (formula (1))

 R_1 , R_2 , R_3 and/or R_4 , is independently from each other selected from H, halogen, hydroxy, protected hydroxy, cyano, nitro, C_1 to C_6 alkyl, C_1 to C_6 substituted alkyl, C_1 to C_7 alkoxy, C_1 to C_7 substituted alkoxy, C_1 to C_7 acyl, C_1 to C_7 substituted acyl, C_1 to C_7 acyloxy, carboxy, protected carboxy, carboxymethyl, protected carboxymethyl, hydroxymethyl, protected hydroxymethyl, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino, carboxamide, protected carboxamide, N-(C_1 to C_6 alkyl)carboxamide, N, N-di(C_1 to C_6 alkyl)carboxamide, N, N-di(C_1 to C_6 al-

kyl)carboxamide, trifluoromethyl, N-((C₁ to C₆ alkyl)sulfonyl)amino, N- (phenylsulfonyl)amino or phenyl, wherein the phenyl is substituted or unsubstituted, such that, for example, a biphenyl results. R₅ is H, C₁ to C₈ alkyl, C₁ to C₈ substituted alkyl, C₇ to C₁₂ alkylphenyl or C₇ to C₁₂ substituted phenylalkyl, R₆ is H, C₁ to C₈ alkyl, C₁ to C₈ substituted alkyl, C₇ to C₁₂ alkylphenyl or C₇ to C₁₂ substituted phenylalkyl, R₇ is H, C₁ to C₈ alkyl, C₁ to C₈ substituted alkyl, C₇ to C₁₂ alkylphenyl or C₇ to C₁₂ substituted phenylalkyl, R₆ and R₇ may be taken together with nitrogen to form a heterocycle or substituted heterocycle or a heteroaryl or substituted heteroaryl ring.

The compounds of the invention can also exist as solvates and hydrates. Thus, these compounds may crystallize with, for example, waters of hydration, or one, a number of, or any fraction thereof of molecules of the mother liquor solvent. The solvates and hydrates of such compounds are included within the scope of this invention.

The symbol "H" denotes a hydrogen atom.

The term " C_1 to C_7 acyl" encompasses groups such as formyl, acetyl, propionyl, butyryl, pentanoyl, pivaloyl, hexanoyl, heptanoyl, benzoyl and the like. Preferred acyl groups are acetyl and benzoyl.

The term "C₁ to C₇ substituted acyl" denotes the acyl group substituted by one or more, and preferably one or two, halogen, hydroxy, protected hydroxy, oxo, protected oxo, cyclohexyl, naphthyl, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino, guanidino, heterocyclic ring, substituted heterocyclic ring, imidazolyl, indolyl, pyrrolidinyl, C₁ to C₇ alkoxy, C₁ to C₇ acyl, C₁ to C₇ acyloxy, nitro, C₁ to C₆ alkyl ester, carboxy, protected carboxy, carbamoyl, carboxamide, protected carboxamide, N-(C₁ to C₆ alkyl)carboxamide, protected N-(C₁ to C₆ alkyl)carboxamide, N,N-di(C₁ to C₆ alkyl)carboxamide, cyano, methylsulfonylamino, thiol, C₁ to C₄ alkylthio or C₁ to C₄ alkylsulfonyl groups. The substituted acyl groups may be substituted once or more, and preferably once or twice, with the same or with different substituents.

The term "substituted phenyl" specifies a phenyl group substituted with one or more, and preferably one or two, moieties chosen from the groups consisting of halogen, hydroxy, protected hydroxy, cyano, nitro, C₁ to C₆ alkyl, C₁ to C₆ substituted alkyl, C₁ to C₇ alkoxy, C₁ to

 C_7 substituted alkoxy, C_1 to C_7 acyl, C_1 to C_7 substituted acyl, C_1 to C_7 acyloxy, carboxy, protected carboxy, carboxymethyl, protected hydroxymethyl, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino, carboxamide, protected carboxamide, N-(C_1 to C_6 alkyl)carboxamide, protected N-(C_1 to C_6 alkyl)carboxamide, N-(N-di(N-di)carboxamide, trifluoromethyl, N-((N-di)carboxamide, N-di)carboxamide, N-di)carboxamide, trifluoromethyl, N-((N-di)carboxamide, N-di)carboxamide, N-di)carboxamide, trifluoromethyl, N-((N-di)carboxamide, N-di)carboxamide, N-di)carboxamide, trifluoromethyl, N-((N-di)carboxamide, N-di)carboxamide, N-di)carboxamide,

Examples of the term "substituted phenyl" includes a mono- or di(halo)phenyl group such as 2, 3 or 4-chlorophenyl, 2,6-dichlorophenyl, 2,5-dichlorophenyl, 3,4-dichlorophenyl, 2, 3 or 4bromophenyl, 3,4-dibromophenyl, 3-chloro-4-fluorophenyl, 2, 3 or 4-fluorophenyl and the like; a mono or di(hydroxy)phenyl group such as 2, 3 or 4-hydroxyphenyl, 2,4-dihydroxyphenyl, the protected-hydroxy derivatives thereof and the like; a nitrophenyl group such as 2, 3 or 4-nitrophenyl; a cyanophenyl group, for example, 2, 3 or 4cyanophenyl; a mono- or di(alkyl)phenyl group such as 2, 3 or 4-methylphenyl, 2,4dimethylphenyl, 2, 3 or 4-(iso-propyl)phenyl, 2, 3 or 4-ethylphenyl, 2, 3 or 4-(npropyl)phenyl and the like; a mono or di(alkoxyl)phenyl group, for example, 2,6-dimethoxyphenyl, 2, 3 or 4-methoxyphenyl, 2, 3 or 4-ethoxyphenyl, 2, 3 or 4-(isopropoxy)phenyl, 2, 3 or 4-(t-butoxy)phenyl, 3-ethoxy-4-methoxyphenyl and the like; 2, 3 or 4-trifluoromethylphenyl; a mono- or dicarboxyphenyl or (protected carboxy)phenyl group such as 2, 3 or 4-carboxyphenyl or 2,4-di(protected carboxy)phenyl; a mono-or di(hydroxymethyl)phenyl or (protected hydroxymethyl)phenyl such as 2, 3, or 4-(protected hydroxymethyl)phenyl or 3,4-di(hydroxymethyl)phenyl; a mono- or di(aminomethyl)phenyl or (protected aminomethyl)phenyl such as 2, 3 or 4-(aminomethyl)phenyl or 2,4-(protected aminomethyl)phenyl; or a mono- or di(N-(methylsulfonylamino))phenyl such as 2, 3 or 4-(N-(methylsulfonylamino))phenyl. Also, the term "substituted phenyl" represents disubstituted phenyl groups wherein the substituents are different, for example, 3-methyl-4hydroxyphenyl. 3-chloro-4-hydroxyphenyl, 2-methoxy-4-bromophenyl. 4-ethyl-2-hydroxyphenyl, 3-hydroxy-4-nitrophenyl, 2-hydroxy 4-chlorophenyl and the like.

The term "heteroaryl" means a heterocyclic aromatic derivative which is a five-membered or six-membered ring system having from 1 to 4 heteroatoms, such as oxygen, sulfur and/or nitrogen, in particular nitrogen, either alone or in conjunction with sulfur or oxygen ring atoms.

Examples of heteroaryls include pyridinyl, pyrimidinyl, and pyrazinyl, pyridazinyl, pyrrolo, furano, thiopheno, oxazolo, isoxazolo, phthalimido, thiazolo and the like.

The term "substituted heteroaryl" means the above-described heteroaryl is substituted with, for example, one or more, and preferably one or two, substituents which are the same or different which substituents can be halogen, hydroxy, protected hydroxy, cyano, nitro, C₁ to C₆ alkyl, C₁ to C₇ alkoxy, C₁ to C₇ substituted alkoxy, C₁ to C₇ acyl, C₁ to C₇ substituted acyl, C₁ to C₇ acyloxy, carboxy, protected carboxy, carboxymethyl, protected carboxymethyl, hydroxymethyl, protected hydroxymethyl, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino, carboxamide, protected carboxamide, N-(C₁ to C₆ alkyl)carboxamide, protected N-(C₁ to C₆ alkyl)carboxamide, N, N-di(C₁ to C₆ alkyl)carboxamide, trifluoromethyl, N-((C₁ to C₆ alkyl)sulfonyl)amino or N-(phenylsulfonyl)amino groups.

The term "substituted naphthyl" specifies a naphthyl group substituted with one or more, and preferably one or two, moieties either on the same ring or on different rings chosen from the groups consisting of halogen, hydroxy, protected hydroxy, cyano, nitro, C₁ to C₆ alkyl, C₁ to C₇ alkoxy, C₁ to C₇ acyl, C₁ to C₇ acyloxy, carboxy, protected carboxy, carboxymethyl, protected carboxymethyl, hydroxymethyl, protected hydroxymethyl, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino, carboxamide, protected carboxamide, N-(C₁ to C₆ alkyl)carboxamide, protected N-(C₁ to C₆ alkyl)carboxamide, trifluoromethyl, N-((C₁ to C₆ alkyl)sulfonyl)amino or N-(phenylsulfonyl)amino.

Examples of the term "substituted naphthyl" includes a mono or di(halo)naphthyl group such as 1, 2, 3, 4, 5, 6, 7 or 8-chloronaphthyl, 2, 6-dichloronaphthyl, 2, 5-dichloronaphthyl, 3, 4-dichloronaphthyl, 1, 2, 3, 4, 5, 6, 7 or 8-bromonaphthyl, 3, 4-dibromonaphthyl, 3-chloro-4-fluoronaphthyl, 1, 2, 3, 4, 5, 6, 7 or 8-fluoronaphthyl and the like; a mono or di(hydroxy)naphthyl group such as 1, 2, 3, 4, 5, 6, 7 or 8-hydroxynaphthyl, 2, 4-dihydroxynaphthyl, the protected-hydroxy derivatives thereof and the like; a nitronaphthyl group such as 3- or 4-nitronaphthyl; a cyanonaphthyl group, for example, 1, 2, 3, 4, 5, 6, 7 or 8-cyanonaphthyl, 1, 2, 4-dimethylnaphthyl, 1, 2, 3, 4, 5, 6, 7 or 8-(isopropyl)naphthyl, 1, 2, 3, 4, 5, 6, 7 or 8-ethylnaphthylnaphthylnaphthylnaphthylnaphthylnaphthylnaphthylnaphthylnaphthylnaphthylnaphthylnaphthylnaphth

di(alkoxy)naphthyl group, for example, 2, 6-dimethoxynaphthyl, 1, 2, 3, 4, 5, 6, 7 or 8-methoxynaphthyl, 1, 2, 3, 4, 5, 6, 7 or 8-ethoxynaphthyl, 1, 2, 3, 4, 5, 6, 7 or 8-(isopropoxy)naphthyl, 1, 2, 3, 4, 5, 6, 7 or 8-(t-butoxy)naphthyl, 3-ethoxy-4methoxynaphthyl and the like; 1, 2, 3, 4, 5, 6, 7 or 8-trifluoromethylnaphthyl; a mono- or dicarboxynaphthyl or (protected carboxy)naphthyl group such as 1, 2, 3, 4, 5, 6, 7 or 8carboxynaphthyl or 2, 4-di(-protected carboxy)naphthyl; a mono-or di(hydroxymethyl)naphthyl or (protected hydroxymethyl)naphthyl such as 1, 2, 3, 4, 5, 6, 7 or 8-(protected hydroxymethyl)naphthyl or 3, 4-di(hydroxymethyl)naphthyl; a mono- or di(amino)naphthyl or (protected amino)naphthyl such as 1, 2, 3, 4, 5, 6, 7 or 8-(amino)naphthyl or 2, 4-(protected amino)-naphthyl, a mono- or di(aminomethyl)naphthyl or (protected aminomethyl)naphthyl such as 2, 3, or 4-(aminomethyl)naphthyl or 2, 4-(protected aminomethyl)-naphthyl; or a mono- or di-(N-methylsulfonylamino) naphthyl such as 1, 2, 3, 4, 5, 6, 7 or 8-(N-methylsulfonylamino)naphthyl. Also, the term "substituted naphthyl" represents disubstituted naphthyl groups wherein the substituents are different, for example, 3methyl-4-hydroxynaphth-1-yl, 3-chloro-4-hydroxynaphth-2-yl, 2-methoxy-4-bromonaphth-1yl, 4-ethyl-2-hydroxynaphth-1-yl, 3-hydroxy-4-nitronaphth-2-yl, 2-hydroxy-4-chloronaphth-1-yl, 2-methoxy-7-bromonaphth-1-yl, 4-ethyl-5-hydroxynaphth-2-yl, 3-hydroxy-8nitronaphth-2-yl, 2-hydroxy-5-chloronaphth-1-yl and the like.

The term "C₁ to C₈ alkyl" denotes such radicals as methyl, ethyl, n-propyl, isopropyl, n-butyl, iso-butyl, sec-butyl, tert-butyl, amyl, tert-amyl, hexyl, n-heptyl, 2-heptyl, 3-heptyl, 4-heptyl, 2-methyl-1hexyl, 2-methyl-3-hexyl, n-octyl and the like.

Examples of the above substituted alkyl groups include the 2-oxo-prop-1-yl, 3-oxo-but-1-yl, cyanomethyl, nitromethyl, chloromethyl, hydroxymethyl, tetrahydropyranyloxymethyl, trityloxymethyl, propionyloxymethyl, amino, methylamino, aminomethyl, dimethylamino, carboxymethyl, allyloxycarbonylmethyl, allyloxycarbonylaminomethyl, methoxymethyl, ethoxymethyl, t-butoxymethyl, acetoxymethyl, chloromethyl, bromomethyl, iodomethyl, trifluoromethyl, 6-hydroxyhexyl, 2,4-dichloro(n-butyl), 2-aminopropyl, 1-chloroethyl, 2-chloroethyl, 1-fluoroethyl, 2-fluoroethyl, 1- iodoethyl, 2-iodoethyl, 1-chloropropyl, 2-chloropropyl, 3- chloropropyl, 1-bromopropyl, 2-bromopropyl, 3-bromopropyl, 1-fluoropropyl, 2-fluoropropyl, 3-fluoropropyl, 1- iodopropyl, 2-iodopropyl, 3-iodopropyl, 2-aminoethyl, N-acetyl-2-aminoethyl, N-acetyl-1-aminoethyl, N-acetyl-1-aminoethyl, and the like.

The term "C₁ to C₈ substituted alkyl" denotes that the above C₁ to C₈ alkyl groups are substituted by one or more, and preferably one or two, halogen, hydroxy, protected hydroxy, oxo, protected oxo, C₃ to C₇ cycloalkyl, naphthyl, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino, guanidino, protected guanidino, heterocyclic ring, substituted heterocyclic ring, imidazolyl, indolyl, pyrrolidinyl, C₁ to C₇ alkoxy, C₁ to C₇ acyl, C₁ to C₇ acyloxy, nitro, carboxy, protected carboxy, carbamoyl, carboxamide, protected carboxamide, N-(C₁ to C₆ alkyl)carboxamide, protected N-(C₁ to C₆ alkyl)carboxamide, N,N-di(C₁ to C₆ alkyl)carboxamide, cyano, methylsulfonylamino, thiol, C₁ to C₄ alkylthio or C₁ to C₄ alkylsulfonyl groups. The substituted alkyl groups may be substituted once or more, and preferably once or twice, with the same or with different substituents.

The term " C_7 to C_{12} phenylalkyl" denotes a C_1 to C_6 alkyl group substituted at any position by a phenyl, substituted phenyl, heteroaryl or substituted heteroaryl. Examples of such a group include benzyl, 2-phenylethyl, 3-phenyl(n-propyl), 4-phenylhexyl, 3-phenyl(n-amyl), 3-phenyl(sec-butyl) and the like. Preferred C_7 to C_{12} phenylalkyl groups are the benzyl and the phenylethyl groups.

The term "C₇ to C₁₂ substituted phenylalkyl" denotes a C₇ to C₁₂ phenylalkyl group substituted on the C₁ to C₆ alkyl portion with one or more, and preferably one or two, groups chosen from halogen, hydroxy, protected hydroxy, oxo, protected oxo, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino, guanidino, protected guanidino, heterocyclic ring, substituted heterocyclic ring, C₁ to C₆ alkyl, C₁ to C₆ substituted alkyl, C₁ to C₇ alkoxy, C₁ to C₇ substituted alkoxy, C₁ to C₇ acyl, C₁ to C₇ substituted acyl, C₁ to C₇ acyloxy, nitro, carboxy, protected carboxy, carbamoyl, carboxamide, protected carboxamide, N-(C₁ to C₆ alkyl)carboxamide, protected N-(C₁ to C₆ alkyl)carboxamide, N, N-(C₁ to C₆ dialkyl)carboxamide, cyano, N-(C₁ to C₆ alkylsulfonyl)amino, thiol, C₁ to C₄ alkylthio, C₁ to C₄ alkylsulfonyl groups; and/or the phenyl group may be substituted with one or more, and preferably one or two, substituents chosen from halogen, hydroxy, protected hydroxy, cyano, nitro, C₁ to C₆ alkyl, C₁ to C₆ substituted alkyl, C₁ to C₇ alkoxy, C₁ to C₇ substituted alkoxy, C₁ to C₇ acyloxy, carboxy, protected carboxy, carboxymethyl, protected carboxymethyl, hydroxymethyl, protected hydroxymethyl, amino, protected amino, (monosubstituted)amino,

protected (monosubstituted)amino, (disubstituted)amino, carboxamide, protected carboxamide, N-(C_1 to C_6 alkyl) carboxamide, protected N-(C_1 to C_6 alkyl) carboxamide, N, N-di(C_1 to C_6 alkyl)carboxamide, trifluoromethyl, N-((C_1 to C_6 alkyl)sulfonyl)amino, N-(phenylsulfonyl)amino, cyclic C_2 to C_7 alkylene or a phenyl group, substituted or unsubstituted, for a resulting biphenyl group. The substituted alkyl or phenyl groups may be substituted with one or more, and preferably one or two, substituents which can be the same or different.

Examples of the term "C₇ to C₁₂ substituted phenylalkyl" include groups such as 2-phenyl-1-chloroethyl, 2-(4-methoxyphenyl)ethyl, 4-(2,6-dihydroxy phenyl)n-hexyl, 2-(5-cyano-3-methoxyphenyl)n-pentyl, 3-(2,6-dimethylphenyl)n-propyl, 4-chloro-3-aminobenzyl, 6-(4-methoxyphenyl)-3-carboxy(n-hexyl), 5-(4-aminomethylphenyl)- 3-(aminomethyl)n-pentyl, 5-phenyl-3-oxo-n-pent-1-yl and the like.

As outlined above R₆ and R₇ may be taken together with nitrogen to form a heterocycle or substituted heterocycle of the following kind aziridine, azetidine, pyrrolidine, 3-methylpyrrolidine, 3-aminopyrrolidine, 3-hydroxypyrrolidine, pyrazolidine, imidazolidine, piperidine, 2-methylpiperidine, piperazine, morpholine, azepine, tetrahydroisoquinoline

The term "heterocycle" or "heterocyclic ring" denotes optionally substituted five-membered to eight-membered rings that have 1 to 4 heteroatoms, such as oxygen, sulfur and/or nitrogen, in particular nitrogen, either alone or in conjunction with sulfur or oxygen ring atoms. These five-membered to eight-membered rings may be saturated, fully unsaturated or partially unsaturated, with fully saturated rings being preferred. Preferred heterocyclic rings include morpholino, piperidinyl, piperazinyl, 2-amino-imidazoyl, tetrahydrofurano, pyrrolo, tetrahydrothiophen-yl, hexylmethyleneimino and heptylmethyleneimino.

The term "substituted heterocycle" or "substituted heterocyclic ring" means the above-described heterocyclic ring is substituted with, for example, one or more, and preferably one or two, substituents which are the same or different which substituents can be halogen, hydroxy, protected hydroxy, cyano, nitro, C₁ to C₁₂ alkyl, C₁ to C₁₂ alkoxy, C₁ to C₁₂ substituted alkoxy, C₁ to C₁₂ acyl, C₁ to C₁₂ acyloxy, carboxy, protected carboxy, carboxymethyl, protected carboxymethyl, hydroxymethyl, protected hydroxymethyl, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino carbox-

amide, protected carboxamide, N-(C_1 to C_{12} alkyl)carboxamide, protected N-(C_1 to C_{12} alkyl)carboxamide, N, N-di(C_1 to C_{12} alkyl)carboxamide, trifluoromethyl, N-((C_1 to C_{12} alkyl)sulfonyl)amino, N-(phenylsulfonyl)amino, heterocycle or substituted heterocycle groups.

The term " C_1 to C_8 alkoxy" as used herein denotes groups such as methoxy, ethoxy, n-propoxy, isopropoxy, n-butoxy, t-butoxy and like groups. A preferred alkoxy is methoxy. The term " C_1 to C_8 substituted alkoxy" means the alkyl portion of the alkoxy can be substituted in the same manner as in relation to C_1 to C_8 substituted alkyl.

The term "C₁ to C₈ aminoacyl" encompasses groups such as formyl, acetyl, propionyl, butyryl, pentanoyl, pivaloyl, hexanoyl, heptanoyl, octanoyl, benzoyl and the like.

The term "C₁ to C₈ substituted aminoacyl" denotes the acyl group substituted by one or more, and preferably one or two, halogen, hydroxy, protected hydroxy, oxo, protected oxo, cyclohexyl, naphthyl, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino, guanidino, heterocyclic ring, substituted heterocyclic ring, imidazolyl, indolyl, pyrrolidinyl, C₁ to C₁₂ alkoxy, C₁ to C₁₂ acyl, C₁ to C₁₂ acyloxy, nitro, C₁ to C₁₂ alkyl ester, carboxy, protected carboxy, carbamoyl, carboxamide, protected carboxamide, N-(C₁ to C₁₂ alkyl)carboxamide, protected N-(C₁ to C₁₂ alkyl)carboxamide, N,N-di(C₁ to C₁₂ alkyl)carboxamide, cyano, methylsulfonylamino, thiol, C₁ to C₁₀ alkylthio or C₁ to C₁₀ alkylsulfonyl groups. The substituted acyl groups may be substituted once or more, and preferably once or twice, with the same or with different substituents.

Examples of C_1 to C_8 substituted acyl groups include 4-phenylbutyroyl, 3-phenylpropanoyl, 2- cyclohexanylacetyl, cyclohexanecarbonyl, 2-furanoyl and 3-dimethylaminobenzoyl.

This invention provides a pharmaceutical composition comprising an effective amount of a compound according to the invention. Such compounds can be administered by various routes, for example oral, subcutaneous, intramuscular, intravenous or intracerebral. The preferred route of administration would be oral at daily doses of the compound for adult human treatment of about 0.01 -5000 mg, preferably 1-1500 mg per day. The appropriate dose may be administered in a single dose or as divided doses presented at appropriate intervals for example as two, three four or more subdoses per day.

For preparing pharmaceutical compositions containing compounds of the invention, inert, pharmaceutically acceptable carriers are used. The pharmaceutical carrier can be either solid or liquid. Solid form preparations include, for example, powders, tablets, dispersible granules, capsules, cachets, and suppositories.

A solid carrier can be one or more substances which can also act as diluents, flavoring agents, solubilizers, lubricants, suspending agents, binders, or tablet disintegrating agents; it can also be an encapsulating material.

In powders, the carrier is generally a finely divided solid which is in a mixture with the finely divided active component. In tablets, the active compound is mixed with the carrier having the necessary binding properties in suitable proportions and compacted in the shape and size desired.

For preparing pharmaceutical composition in the form of suppositories, a low-melting wax such as a mixture of fatty acid glycerides and cocoa butter is first melted and the active ingredient is dispersed therein by, for example, stirring. The molten homogeneous mixture is then poured into convenient-sized molds and allowed to cool and solidify.

Powders and tablets preferably contain between about 5% to about 70% by weight of the active ingredient. Suitable carriers include, for example, magnesium carbonate, magnesium stearate, talc, lactose, sugar, pectin, dextrin, starch, tragacanth, methyl cellulose, sodium carboxymethyl cellulose, a low-melting wax, cocoa butter and the like.

The pharmaceutical compositions can include the formulation of the active compound with encapsulating material as a carrier providing a capsule in which the active component (with or without other carriers) is surrounded by a carrier, which is thus in association with it. In a similar manner, cachets are also included. Tablets, powders, cachets, and capsules can be used as solid dosage forms suitable for oral administration.

Liquid pharmaceutical compositions include, for example, solutions suitable for oral or parenteral administration, or suspensions, and emulsions suitable for oral administration. Sterile water solutions of the active component or sterile solutions of the active component in solvents comprising water, ethanol, or propylene glycol are examples of liquid compositions suitable for parenteral administration.

Sterile solutions can be prepared by dissolving the active component in the desired solvent system, and then passing the resulting solution through a membrane filter to sterilize it or, alternatively, by dissolving the sterile compound in a previously sterilized solvent under sterile conditions.

In one embodiment of the present invention a compound is claimed according to formula (1) above, or pharmaceutical acceptable salts or solvates thereof, wherein R₁, R₂, R₃, R₄, is H, halogen, hydroxy, protected hydroxy, cyano, nitro, C₁ to C₆ alkyl, C₁ to C₆ substituted alkyl, C₁ to C₇ alkoxy, C₁ to C₇ substituted alkoxy, C₁ to C₇ acyl, C₁ to C₇ substituted acyl, C₁ to C₇ acyloxy, carboxy, protected carboxy, carboxymethyl, protected carboxymethyl, protected hydroxymethyl, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino, carboxamide, protected carboxamide, N-(C₁ to C₆ alkyl)carboxamide, protected N-(C₁ to C₆ alkyl)carboxamide, N, N-di(C₁ to C₆ alkyl)carboxamide, trifluoromethyl, N-((C₁ to C₆ alkyl)sulfonyl)amino, N- (phenylsulfonyl)amino or phenyl, R₅ is H, C₁ to C₈ alkyl, C₁ to C₈ substituted alkyl, C₇ to C₁₂ alkylphenyl or C₇ to C₁₂ substituted phenylalkyl, R₆ and R₇ may be taken together with nitrogen to form the heterocycle according to formula (2),

formula (2)



In a preferred embodiment of the invention a compound is provided, or pharmaceutical acceptable salts or solvates thereof, wherein R₁, R₂, R₃, R₄, is H, halogen, hydroxy, protected hydroxy, cyano, nitro, C₁ to C₆ alkyl, C₁ to C₆ substituted alkyl, C₁ to C₇ alkoxy, C₁ to C₇ substituted alkoxy, C₁ to C₇ acyl, C₁ to C₇ substituted acyl, C₁ to C₇ acyloxy, carboxy, protected carboxy, carboxymethyl, protected carboxymethyl, hydroxymethyl, protected hydroxymethyl, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino, carboxamide, protected carboxamide, N-(C₁ to C₆ alkyl)carboxamide, pro-

tected N-(C_1 to C_6 alkyl)carboxamide, N, N-di(C_1 to C_6 alkyl)carboxamide, trifluoromethyl, N-((C_1 to C_6 alkyl)sulfonyl)amino, N- (phenylsulfonyl)amino or phenyl, R_5 is H, C_1 to C_8 alkyl, C_1 to C_8 substituted alkyl, R_6 and R_7 may be taken together with nitrogen to form the heterocycle according to formula (2) shown above.

A particularly preferred compound which may act as agonist of LXR is shown in formula (6) below. The inventors have been able to demonstrate that the compound according to formula (3) has a low effective concentration at LXR with an EC₅₀ of 0.5 μ M wherein the EC₅₀ reflects the half-maximal effective concentration, and which is higher than the EC₅₀ of 0.015 μ M for the published LXR agonist TO901317 (J. Schultz et al., Genes Dev. 14, 2831-2838, 2000)

formula (3) (MOLNAME 3252)

The inventors have also found the compounds according to formula (4, 5 and 6) (shown below) to be active as agonist of the LXR human nuclear receptor (see figures for details).

formula (4) (MOLNAME 7459)

formula (5) (MOLNAME 6584)

formula (6)) (MOLNAME 7364)

In particular the invention relates to a compound as described above wherein said compounds is capable of binding the LXR receptor protein or a portion thereof according to SEQ ID NO. 1 (Fig. 3 A to F) or a mammalian homologue thereof. The claimed compound can bind to the LXR receptor protein or a portion thereof in a mixture comprising 10-200 ng of LXR receptor protein, a fusion protein containing LXR or a portion thereof, preferably the ligand binding domain, fused to a Tag, 5-100 mM Tris /HCl at pH 6,8-8,3; 60-1000 mM KCl; 0-20 mM MgCl2; 100-1000ng/μl BSA in a total volume of preferably about 25 μl.).

A mammalian receptor protein homologue of the protein according to SEQ ID NO. 1 as used herein is a protein that performs substantially the same task as LXR does in humans and shares at least 40% sequence identity at the amino acid level, preferably over 50 % sequence identity at the amino acid level more preferably over 65 % sequence identity at the amino acid level, even more preferably over 75 % sequence identity at the amino acid level and most preferably over 85 % sequence identity at the amino acid level.

The invention in particular concerns a method for prevention or treatment of a LXR receptor protein or LXR receptor protein homologue mediated disease or condition in a mammal comprising administration of a therapeutically effective amount of a compound according to the invention wherein the prevention or treatment is directly or indirectly accomplished through

the binding of a compound according to the invention to the LXR receptor protein or to the LXR receptor protein homologue.

The term mediated herein means that the physiological pathway in which the LXR receptor protein acts is either directly or indirectly involved in the disease or condition to be treated or prevented. In the case where it is indirectly involved it could be that, e.g. modulating the activity of LXR by a compound according to the invention influences a parameter which has a beneficial effect on a disease or a condition. One such example is that modulation of LXR activity leads to decreased levels of serum cholesterol or certain lipoproteins which in turn have a beneficial effect on the prevention and treatment of atherosclerosis. Herein a condition is a physiological or phenotypic state which is desirably altered. One such example would be obesity which is not necessarily medically harmful but nonetheless a non desirable phenotypic condition. In a preferred embodiment of the invention the method for prevention or treatment of a LXR receptor protein mediated disease or condition is applied to a human. This may be male or female.

Pharmaceutical compositions generally are administered in an amount effective for treatment or prophylaxis of a specific condition or conditions. Initial dosing in human is accompanied by clinical monitoring of symptoms, such symptoms for the selected condition. In general, the compositions are administered in an amount of active agent of at least about 100 μg/kg body weight. In most cases they will be administered in one or more doses in an amount not in excess of about 20 mg/kg body weight per day. Preferably, in most cases, doses is from about 100 μg/kg to about 5 mg/kg body weight, daily.

For administration particularly to mammals, and particularly humans, it is expected that the daily dosage level of active agent will be 0,1 mg/kg to 10 mg/kg and typically around 1 mg/kg.

By "therapeutically effective amount" is meant a symptom- alleviating or symptom -reducing amount, a cholesterol-reducing amount, a cholesterol absorption blocking amount, a protein and/or carbohydrate digestion-blocking amount and/or a de novo cholesterol biosynthesis-blocking amount of a compound according to the invention.

Likewise, the invention concerns a method of treating in mammal a disease which is correlated with abnormal cholesterol, triglyceride, or bile acid levels or deposits comprising administering to a mammal in need of such treatment a therapeutically effective amount of a compound according to the invention.

Accordingly, the compounds according to the invention may also be used as a method of prevention or treatment of mammalian atherosclerosis, gallstone disease, lipid disorders, Alzheimer's disease, skin disorders, obesity or cardiovascular disorders such as coronary heart disease or stroke.

The invention further concerns a method of blocking in a mammal the cholesterol absorption in the intestine in need of such blocking comprising administering to a mammal in need of such treatment a therapeutically effective amount of a compound according to the invention. The invention may also be used to treat obesity in humans.

The Liver X Receptor alpha is a prototypical type 2 nuclear receptor meaning that it activates genes upon binding to the promoter region of target genes in a heterodimeric fashion with Retinoid X Receptor. The relevant physiological ligands of LXR are oxysterols The present compounds according to the invention have been demonstrated to have a high binding efficacy (binding coefficients measured as EC50 in the range 100 nM to 1500 nM) as well as agonistic and / or antagonistic properties. Consequently they may be applied to regulate genes that participate in bile acid, cholesterol and fatty acid homeostasis as well as other downstream regulated genes. Examples of such genes are but are not limited to lipid absorption, cholesterol biosynthesis, cholesterol transport or binding, bile acid transport or binding, proteolysis, amino acid metabolism, glucose biosynthesis, protein translation, electron transport, and hepatic fatty acid metabolism. LXR often functions in vivo as a heterodimer with the Retinoid X Receptor. Published LXR agonists such as the Tularik compound "TO901317" (See figure 5) are known to influence the regulation of various liver genes. Genes found to be regulated by TO901317 can be found in figure 6. Thus, the invention also concerns a method of modulating a gene whose expression is regulated by the LXR receptor in a mammal comprising administration of a therapeutically effective amount of a compound according to the invention to said mammal.

A number of direct and indirect LXR target genes have been described whose regulated expression contribute to cholesterol homeostasis and lipogenesis. In this respect the direct regulation of Cyp7A, which was shown to be a direct target gene of LXR at least in the rodent lineage is an important aspect of cholesterol removal by increased metabolism of bile acids (Lehmann et al., J Biol.Chem. 272 (6) 3137-3140; 1007). Gupta et al. (Biochem. Biophys Res.Com, 293; 338-343, 2002) showed that LXR α regulation of Cyp7A is dominant over FXR inhibitory effects on Cyp7A transcription.

A key transcription factor that was also shown to be a direct target gene for the LXR receptor is SREBP-1C (Repa et al., Genes and Development, 14:2819-2830; 2000: Yoshikawa et al.; Mol.Cell.Biol.21 (9) 2991-3000, 2001). SREBP-1C itself activates transcription of genes involved in cholesterol and fatty acid synthesis in liver but also other mammalian tissues. Some of the SREBP1c target genes involved in lipogenesis like FAS and SCD have shown to be additionally direct targets of the LXR receptors (Joseph et al.; J Biol Chem. 2002 Mar 29;277(13):11019-25; Liang et al., J Biol Chem. 2002 Mar 15;277(11):9520-8.).

Another gene that has been shown to be directly regulated by LXRs is the LPL gene, that codes for a key enzyme that is responsible for the hydrolysis of triglycerides in circulating lipoprotein, releasing free fatty acids to peripheral tissues. (Zhang et al. J Biol Chem. 2001 Nov 16;276(46):43018-24.) This enzyme is believed to promote uptake of HDL cholesterol in liver, thereby promoting reverse cholesterol transport. A similar functional involvement in HDL clearance is described for the CETP gene product that facilitated the transfer of HDL cholesterol esters from plasma to the liver. LXR response elements were found in the CETP promoter and direct activation of this gene by LXR was demonstrated (Luo and Tall; J Clin Invest. 2000 Feb;105(4):513-20.).

The regulated transport of cholesterol through biological membranes is an important mechanism in order to maintain cholesterol homeostasis. A pivotal role in these processes in multiple tissues like e.g. macrophages and intestinal mucosa cells is maintained by the ATP-binding cassette transporter proteins (ABC). ABCA1 and ABCG1 were identified as direct LXR target genes (Costet et al.; J Biol Chem. 2000 Sep 8;275(36):28240-5) that mediate cholesterol efflux and prevent thereby e.g. generation of artherogenic plaques in macrophages (Singaraja et al. J Clin Invest. 2002 Jul;110(1):35-42). Other ABC transporters like ABCG5

and ABCG8, primarily expressed in hepatocytes and enterocytes have also been reported to be directly responsive to LXR agonists (Repa et al., J Biol Chem. 2002 May 24;277(21):18793-800. Kennedy et al., J Biol Chem. 2001 Oct 19;276(42):39438-47) and mediate the secretion of sterols from the liver and efflux of dietary sterols from the gut.

Apolipoproteins E, C-I, C-II, and C-IV, that fulfill important roles in lipoprotein/lipid homeostasis have also been shown to be direct targets of the LXR receptor (Laffitte et al., Proc Natl Acad Sci U S A. 2001 Jan 16;98(2):507-12; Mak et al.; J Biol Chem. 2002 May 24 [epub ahead of print]). These proteins have been found to be crucial components of chylomicrons, VLDL, IDL, and HDL and are among other things associated with hypertriglyceridemia and arteriosclerosis.

Recently the LXRα itself was shown to be regulated by both LXR receptors in human cell types including macrophages suggesting an autoregulatory amplification event in the response to LXR ligands which could e.g. lead to an enhanced stimulation of LXR target genes like e.g. ABCA1 (Bolten et al.; Mol Endocrinol. 2002 Mar;16(3):506-14.; Laffitte et al., Mol Cell Biol. 2001 Nov;21(22):7558-68; Whitney et al.; J Biol Chem. 2001 Nov 23;276(47):43509-15).

Besides the important function of LXR receptors in tissues like liver and macrophages it has recently been reported that that stimulation of epidermal differentiation is mediated by Liver X receptors in murine epidermis. Differentiation maker genes like involucrin, loricin and profilaggrin have been shown to be upregulated upon LXR ligand treatment (Kömüves et al.; J Invest Dermatol. 2002 Jan;118(1):25-34.).

Another recent report describes the regulation of cholesterol homeostasis (primarily the regulation of ABCA1, ABCG1 and SREBP-1C) by the LXR receptors in the central nervous system suggesting that LXRs may prove beneficial in the treatment of CNS diseases such as Alzheimer's and Niemann-Pick disease that are known to be accompanied by dysregulation of cholesterol balance (Whitney et al.; Mol Endocrinol. 2002 Jun;16(6):1378-85).

Therefore one important embodiment the invention concerns are methods that enhances or suppresses amongst other today yet unknown LXR target genes the above mentioned genes and the associated biological processes and pathways through LXR compounds that are subject of this invention.

The compounds according to the invention may be used as medicaments, in particular for the manufacture of a medicament for the prevention or treatment of a LXR receptor protein or LXR receptor protein homologue mediated disease or condition in a mammal wherein the prevention or treatment is directly or indirectly accomplished through the binding of the compound according to the invention to the LXR receptor protein or LXR receptor protein homologue. These pharmaceutical compositions contain 0,1 % to 99,5 % of the compound according to the invention, more particularly 0,5 % to 90 % of the compound according to the invention with a pharmaceutically acceptable carrier.

The invention concerns also the use of a compound according to the invention for the manufacture of a medicament for the prevention or treatment of a LXR receptor protein mediated disease or condition wherein the mammal described above is a human. The medicament may be used for regulating the cholesterol transport system, for regulating levels of cholesterol, triglyceride, and/or bile acid in a mammal preferentially a human by activating the LXR receptor. The medicament may be used for the treatment of atherosclerosis, gallstone disease, lipid disorders, Alzherimer's disease, skin disorders, obesity or a cardiovascular disorder.

The further concerns the use of a compound according to the invention for the manufacture of a medicament capable for blocking in a mammal, preferentially a human the cholesterol absorption in the intestine. Further the claimed compound may be used for the manufacture of a medicament for treating obesity in humans and for modulating a gene whose expression is regulated by the LXR receptor (see details above and figures).

The present invention shall now be further illustrated based on the following examples without being limited thereto. In the accompanying sequence protocol and the figures:

SEQ ID NO. 1 shows protein sequence of the LRX alpha protein a portion of which was used for cloning as described in the examples,

SEQ ID NO. 2 shows the mRNA sequence of the LRX alpha protein,

SEQ ID NO. 3 shows the protein sequence of TIF2 (Acc. No: XM 011633 RefSeq DB),

SEQ ID NO. 4 shows the respective mRNA sequence corresponding to the TIF2 protein,

SEQ ID NO 5 shows the protein sequence of the LXR beta protein a portion of which was used for cloning as described in examples,

SEQ ID NO 6 shows the mRNA sequence of the LXR beta protein,

SEQ ID NO 7 shows the sequence of primer (a) used in Example 1

SEQ ID NO 8 shows the sequence of primer (b) used in Example 1.

Fig. 1 shows the synthesis of the compounds according to the invention as also described in Example 2.

Fig. 2 shows the measurement parameters employed by the Wallace VICTOR2VTM Multilabel Counter which was used for measuring the EC₅₀ values

Fig. 3 A shows SEQ ID NO. 1 which is the protein sequence of the LRX alpha protein a portion of which was used for cloning as described in the examples. Figure 3 B shows SEQ ID NO. 2 which is the mRNA sequence of the LRX alpha protein. Figure 3 C shows SEQ ID NO. 3 which is the protein sequence of TIF2 (Acc. No: XM_011633 RefSeq DB), Figure 3 D shows SEQ ID NO. 4 which is the respective mRNA sequence corresponding to the TIF2 protein. Figure 3 E shows SEQ ID NO 5 which is the protein sequence of the LXR beta protein a portion of which was used for cloning as described in examples. Figure 3 F shows SEQ ID NO 6 which is the mRNA sequence of the LXR beta protein.

Fig. 4 shows the internal molecular name used by the applicant (MOLNAME) as well as the corresponding structures of preferred compounds according to the invention. The figure further shows their respective EC₅₀ values (EC50 AVG) as established according to the experiment 1 in multiple experiments (see above), as well as their respective average efficacy (% activity relative to 22-(R)-hydroxycholesterol control agonist).

Figure 5 shows various known LXR ligands. It is apparent from their structures that the inventors have identified novel compounds which are structurally not related to these known ligands.

Figure 6 shows various genes that have been found to be regulated through binding of an LXR agonist to the LXR protein.

Figure 7 shows a dose-dependent transactivation (EC50 \sim 3 μ M) by LN0000007465 of the luciferase reporter gene via LXR alpha.

Figure 8 shows (A) Analysis of mRNA content of the indicated genes in total RNA isolated from THP-1 cells treated for 24 hours with 2, 10 or 25μ M of LN0000006500 or 10 μ M of the Tularik compound (T0901317). (B) Analysis of mRNA content fo the indicated genes in total RNA from HepG2 cells treated for 24 hours with 2, 10 or 25 μ M of LN0000006500 or 10 μ M of the Tularik compound (T0901317).

Figure 9 shows the dose dependent transactivation by LN0000006500 of the pFR-luc reporter gene in CHO cells via Gal4 LBD-fusion constructs derived from LXRa- or LXRb. Concentrations of the compound administered (μM) and RLU's determined from extracts of cells are indicated.

Figure 10 shows the analysis of total cholesterol from supernatants of cultivated THP-1 cells incubated without or with ApoA1 and ApoA1 plus $10\mu M$ of the compounds Tularik (T0901317) or LN0000006500, LN0000006662, LN00000006671 or LN0000006672 as indicated.

EXAMPLES

EXAMPLE 1:

In vitro screening for compounds which influence LXR binding to coactivators.

For screening purposes a GST and 6 x His fusion of the LBD (from amino acids 155 of hLXRalpha to 447) of human LXRalpha was constructed by first cloning a Gateway cassette (Invitrogen) in frame into the Sma I site of the pAGGHLT Polylinker (Pharmingen). Then a PCR fragment specifically amplified from human liver cDNA was cloned into the resulting

pACGHLT-GW following the manufacturers instructions for Gateway cloning (Invitrogen) to yield pACGHLT-GW-hLXRalphaLBD.

Primers used for amplification were: primer (a) GGGGACAAGTTTGTACAAAAAAGCAGGCTCGCTTCGCAAATGCCGTCAG (SEQ ID NO 7), and primer (b) GGGGACCACTTTGTACAAGAAAGCTGGGTCCCCTTCTCAGTCTGTTCCACTT (SEQ ID NO 8).

100 % sequence integrity of all recombinant products was verified by sequencing. Recombinant Baculovirus was constructed from pACGHLT-GW-hLXRalphaLBD using the Pharmingen Baculovirus Expression vector system according to instructions of the manufacturer. Monolayer cultures of SF9 cells were infected by the virus as recommended by Pharmingen or 200ml cultures of 1 x10⁶ cells/ml grown in 2 liter Erlenmeyer flasks on an orbital shaker at 30 rpm were infected by 10ml of same virus stock. In both cases cells were harvested 3 days after infection. All cell growth was performed in Gibco SF900 II with Glutamine (Invitrogen) medium without serum supplementation at 28°C. Since SF9 cells contain significant amounts of endogenous GST, purification was performed via His and not via GST affinity chromatography. To this end instructions of Pharmingen for purification of recombinant His tagged proteins from SF9 cells were followed with the following modifications: All detergents were omitted from the buffers and cells were lysed on ice by 5 subsequent sonication pulses using a sonicator needle at maximum power.

All eluates were dialyzed against 20 mM Tris/HCl pH 6,8, 300 mM KCl; 5 mM MgCl₂; 1 mM DTT; 0,2 mM PMSF; 10% Glycerol. A typical dialyzed eluate fraction contained the fusion protein at a purity of more than 80%. Total protein concentration was 0,1-0,3 mg/ml.

For E. coli expression of a NR coactivator, pDest17-hTif2BD expressing a NR interaction domain from amino acids 548-878 of human Tif2 (Acc. No: XM_011633 RefSeq) tagged by 6 N-terminal His residues was constructed. Therefore, a PCR fragment specifically amplified from human liver cDNA was subcloned into pDest 17 (Invitrogen) following the manufacturers instructions for Gateway cloning (Invitrogen). Primers used for Amplification were: primer (a)

GGGGACAAGTTTGTACAAAAAAGCAGGCTCGTTAGGGTCATCGTTGGCTTCACC

and primer (b)

GGGGACCACTTTGTACAAGAAAGCTGGGTCTCAAAGTTGCCCTGGTCGTGGGTTA

For E. coli expression plasmid DNA was transformed into chemically competent E. coli BL21 (Invitrogen, USA) and cells were grown to an OD600 of 0.4-0.7 before expression was induced by addition of 0,5 mM IPTG according instructions of the manufacturer (Invitrogen). After induction for 8 hours at 30°C cells were harvested by centrifugation for 10 minutes at 5000 x g. Fusion proteins were affinity purified using Ni-NTA Agarose (QIAGEN) according to the instructions of the manufacturer. Recombinant Tif2 construct was dialyzed against 20 mM Tris/HCL pH 7.9; 60 mM KCl; 5 mM MgCl₂; 1 mM DTT, 0,2 mM PMSF; 10% glycerol. A typical dialyzed eluate fraction contained the fusion protein at a purity of more than 80%. Total protein concentration was 0,1-0,3 mg/ml.

The TIF2 fragment was subsequently biotinylated by addition of 5-40µl/ml Tif2 fraction of a Biotinamidocaproate N-Hydroxysuccinimide-ester (Sigma) solution (20 mg/ml in DMSO). Overhead rotating samples were incubated for 2 hours at room temperature. Unincorporated label was then separated using G25 Gel filtration chromatography (Pharmacia Biotech, Sweden). Protein containing fractions from the column were pooled and tested for activity in the assay as described below.

For screening of compound libraries as provided for by the methods shown below in the examples for substances which influence the LXR/Tif 2 interaction, the Perkin Elmer LANCE technology was applied. This method relies on the binding dependent energy transfer from a donor to an acceptor fluorophore attached to the binding partners of interest. For ease of handling and reduction of background from compound fluorescence LANCE technology makes use of generic fluorophore labels and time resoved detection (for detailed description see Hemmilä I, Blomberg K and Hurskainen P, Time-resolved resonance energy transfer (TR-FRET) principle in LANCE, Abstract of Papers Presented at the 3 rd Annual Conference of the Society for Biomolecular Screening, Sep., California (1997))

For screening, 20-200 ng of biotinylated Tif 2 fragment and 10-200 ng of GST-LXR fragment were combined with 0.5-2 nM LANCE Eu-(W1024) labelled anti-GST antibody (Perkin Elmer) and 0,1-0,5µg of highly fluorescent APC-labelled streptavidin (Perkin Elmer,

AD0059) in the presence of 50µM of individual compounds to be screened in a total volume of 25 µl of 20 mM Tris /HCl pH 6,8; 300 mM KCl; 5 mM MgCl2; 100-1000 ng/µl/ BSA DMSO content of the samples was kept below 4%. Samples were incubated for a minimum of 60 minutes in the dark at room temperature in FIA-Plates black 384well med. binding (Greiner).

The LANCE signal was detected by a Perkin Elmer VICTOR2V™ Multilabel Counter applying the detection parameters listed in Fig. 2. The results were visualized by plotting the ratio between the emitted light at 665 nm and at 615 nm. For every batch of recombinant proteins amount of proteins, including BSA and labeling reagents giving the most sensitive detection of hits was determined individually by analysis of dose response curves for 22R Hydroxycholesterol and TO 901317

EXAMPLE 2:

Experimental procedure for the preparation of the compounds according to the invention.

o-AZIDOBENZOIC ACID SYNTHESIS (2)

The anthranilic acid (1, 1 eq., 0.5-1 M) was suspended in 6 M HCl, containing enough AcOH (0-20% dependent upon the anthranilic acid) to facilitate dissolution of the anthranilic acid and/or the intermediate diazonium salt, and cooled to 0 °C. NaNO₂ (1.1 eq., 1.3-2.5 M) dissolved in H₂O was added to the anthranilic acid solution at a rate such that the temperature of the reaction solution remained below 5 °C. The resulting homogeneous solution of the diazonium salt was slowly filtered through a sintered glass funnel into a solution of NaN₃ (1.1 eq., 0.7-1.1 M) and NaOAc (12 eq.) in H₂O. The reaction mixture was stirred/shaken for 30-60 min following cessation of vigorous N₂ evolution. Following acidification of the reaction mixture to pH 1 with concentrated HCl, the mixture was cooled to 0 °C to encourage complete precipitation of the o-azidobenzoic acid. The precipitate was collected by filtration and washed with 6 M HCl (2x) and H₂O (2x). The o-azidobenzoic acid product (2) was dried in vacuo (500 mtorr, 30 °C).

ACYLATION OF HYDROXYMETHYL RESIN (4)

To hydroxymethyl resin (1.0 eq., 1.3 mmol/g) and the o-azidobenzoic acid (1, 2.5 eq.) was added DMF (to give 400 mM o-azidobenzoic acid ,1), CsCO₃ (2.0 eq.) and KI (2.0 eq.). Following agitation of the reaction mixture for 36-48 h, the resin-bound o-azidobenzoic acid (4)

was washed with MeOH (2 cycles), CH₂Cl₂ (3 cycles), MeOH (3 cycles), DMF (3 cycles), MeOH (3 cycles) and CH₂Cl₂ (3 cycles), and dried *in vacuo*.

AZA-WITTIG FORMATION (5)

To the resin-bound *o*-azidobenzoic acid (4,1.0 eq.) was added a solution of PPh₃ (THF, 500 mM, 5.0 eq.). After 6 h, the resin was washed with 3 cycles of the following: THF (3 cycles), toluene (3 cycles), CH₂Cl₂ (3 cycles) and hexanes (3 cycles). Followed by drying *in vacuo* to afford resin bound iminophosphorane (5)

CARBODIIMIDE FORMATION (6)

To the resin-bound iminophosphorane (5, 1 eq.) was added isocyanate (9, 5 eq., 450 mM) dissolved in ClCH₂CH₂Cl. The compounds were shaken at ambient temperature for 16 h, washed with 3 cycles of the following: THF (3 cycles), toluene (3 cycles), CH₂Cl₂ (3 cycles) and hexanes (3 cycles), and dried *in vacuo* to afford carbodiimide (6).

GUANIDINE FORMATION / CYCLIZATION

To the carbodiimide functionalized resin (6) was added secondary amine (10, 0.6 eq., 500 mM) dissolved in ClCH₂CH₂Cl. The reaction mixture was heated to 50 °C in an incubator for 12-72 h to afford 2-aminoquinazoline (8).

All of the final products were analyzed by HPLC using mass and an Evaporative Light Scattering Detector (ELSD) detection to determine purity and identity.

One skilled in the art will be able to arrive at the compounds claimed herein making use of said protocol.

EXAMPLE 3:

This example illustrates that a compound according to the invention (experiments shown were done with MOLSTRUCTURE LN 0000007465 (see figures 4 for structural formula)) can mediate transactivation of LXR mediated transcription in HEK293 cells.

HEK293 cells were grown in 48 well plates and co-transfected with the pTRexDest30 (Invitrogen) derivatives pTRexDest30-hLXRa, pTRexDest30-hRXR□ and the pGL2promoter (Promega) derivative pGL2promoter-LXRRE (each 300 ng of plasmid DNA). The full length

human LXR (accession U68233) and the full length human RXRα (accession P19793) were cloned into the pTRexDest30 applying the manufacturer protocols for the GatewayTM system (Invitrogen).

The LXR response elements (LXRRE) were (upper case and underlined) 5'

CoeffTGGTCA etca AGTTCA aggregate a

Ccctt<u>TGGTCActcaAGTTCA</u> agtgatgatagaattcggatcctt<u>TGGTCActcaAGTTCA</u> agtgA 3' (SEQ ID NO. 5) derived from the rat Cyp7a promoter (Laffite et al., 2001, PNAS 98,pp 507). Luciferase reporter activity was measured in triplicates from extracts of cells after incubating cells in culture medium (DMEM [Gibco-BRL] + 10% FCS [PAA laboratories]) for 16 hours (5% CO₂, 37°C) containing 0,5% DMSO (control) or 0,5% DMSO with increasing concentrations of LN0000007465.

A dose-dependent transactivation (EC50 \sim 3 μM) of the reporter gene by LXRa was observed (Fig. 7).

Example 4:

This example shows that described compounds can increase the abundance of mRNA of target genes for the LXR proteins in THP-1 cells treated with TPA.

THP-1 (3×10^5 cells per dish) cells were seeded in 24 well dishes in 3 ml modified RPMI-1640 medium (ATTC, Cat.No. 30-2001) containing 10%FCS (GIBCO) and 100nM TPA and cultivated at 37^{0} C in 5% CO₂ for 48 hours. The medium was then removed and replaced with medium containing 10% charcoal/stripped FBS (Hyclone) and incubated with LN0000006500 at 2, 10 or 25 μ M concentration or Tularik (T0901317) at 10 μ M for 24 hours as indicated in Fig 8A as an example. HepG2 (4,5x10⁵ cells per dish) were seeded in 24 well dishes in 3 ml DMEM Medium containing 10%FCS (GIBCO) and cultivated at 37^{0} C in 5% CO₂ for 48 hours as indicated in Fig 8B.

After incubation for 24 hours in presence of compound, total RNA was isolated from the cells using a Quiagen RNAeasy kit (Quiagen) according to the manufacturers protocol. The RNA was then reverse transcribed and analyzed by TaqMan Analysis using kits and equipment from Perkin-Elmer known to those knowledgeable in the field.

The fold change of mRNA abundancy of compound treated versus DMSO treated as a control is shown in Figure 8A and B for several analyzed target genes indicated in Figure 8A and B.

Example 5:

This example shows that described compounds can selectively enhance transcription mediated by the LBD's of the respective nuclear receptors LXRa and LXRb.

CHO cells (1x10⁵ cells 96well plate) were co-transfected (Lipofectamine 2000 GIBCO) with pFR-luc (Stratagene) as a reporter gene construct and pCMV-AD derivatives containing the LXRa or LXRb ligand binding domains, which were cloned via the gateway system (GIBCO) described in Example 1, in order to express Gal4DBD-LXRa or Gal4DBD-LXRb fusion proteins.

Cells were grown in DMEM containing 10%FCS at 37^0C in 5% humidified CO2 for 16h in presence of 0,05% DMSO vehicle or 0,032 to $50~\mu$ M LN0000006500 in vehicle (as indicated in Fig.9). Luciferase activity was determined from aliquots of extracts prepared from cells following standard luciferase assay kits and protocols from Promega.

Example 6:

This example shows that described compounds at 10 μ M concentration for 24 hours can increase the reverse cholesterol transport in THP-1 cells that were treated with TPA.

THP-1 (1x10⁶ cells per dish) cells were seeded in 6 well dishes in 3ml modified RPMI1640 medium (ATTC, Cat.No. 30-2001) containing 10%FCS (GIBCO) and 100nM TPA and cultivated at 37^{0} C in 5% CO₂ for 72 hours. The medium was then removed and replaced with fresh medium containing 100 nM TPA and 0,15 % BSA. After 24 h incubation the cells were washed in PBS and 1,5 ml of fresh medium containing either 0,1% DMSO alone or 0,1% DMSO together with $40\mu g/ml$ ApoA1 (Calbiochem) or $40\mu g/ml$ ApoA1 plus the in Fig.10 as an example indicated compounds Tularik (T0901317), LN00000006500, LN0000006662, LN0000006671, LN0000006674 at 10 μ M.

After incubation for 24 hours, total cholesterol was determined from cell supernatant in each of the wells using an enzymatic assay with fluorescence read-out for the determination of cholesterol (Amplex Red Cholesterol Assay Kit (A-12213). The fluorescence readout per mg

WO 2004/024162 PCT/EP2003/007067

of total protein content as determined from cells that were present in the respective well are shown in Figure 9 as an example.

Claims:

1. A compound of the formula (1), or pharmaceutical acceptable salts or solvates thereof according to formula (1)

$$R_{2}$$

$$R_{3}$$

$$R_{4}$$

$$R_{5}$$

$$R_{7}$$

$$R_{6}$$

wherein:

R₁, R₂, R₃ and/or R₄, is independently from each other H, halogen, hydroxy, protected hydroxy, cyano, nitro, C₁ to C₆ alkyl, C₁ to C₆ substituted alkyl, C₁ to C₇ alkoxy, C₁ to C₇ substituted alkoxy, C₁ to C₇ acyl, C₁ to C₇ substituted acyl, C₁ to C₇ acyloxy, carboxy, protected carboxy, carboxymethyl, protected carboxymethyl, hydroxymethyl, protected hydroxymethyl, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino, carboxamide, protected carboxamide, N-(C₁ to C₆ alkyl)carboxamide, protected N-(C₁ to C₆ alkyl)carboxamide, N, N-di(C₁ to C₆ alkyl)carboxamide, trifluoromethyl, N-((C₁ to C₆ alkyl)sulfonyl)amino, N-(phenylsulfonyl)amino or phenyl, wherein the phenyl is substituted or unsubstituted, R₅ is H, C₁ to C₈ alkyl, C₁ to C₈ substituted alkyl, C₇ to C₁₂ alkylphenyl or C₇ to C₁₂ substituted phenylalkyl,

 R_6 is H, C_1 to C_8 alkyl, C_1 to C_8 substituted alkyl, C_7 to C_{12} alkylphenyl or C_7 to C_{12} substituted phenylalkyl, and

 R_7 is H, C_1 to C_8 alkyl, C_1 to C_8 substituted alkyl, C_7 to C_{12} alkylphenyl or C_7 to C_{12} substituted phenylalkyl.

2. A compound according to claim 1 wherein R₆ and R₇ are taken together with nitrogen to form a heterocycle or substituted heterocycle or a heteroaryl or substituted heteroaryl according to the following formula (2).



3. A compound according to claim 2, or pharmaceutical acceptable salts or solvates thereof, wherein:

R₁, R₂, R₃, R₄, is H, halogen, hydroxy, protected hydroxy, cyano, nitro, C₁ to C₆ alkyl, C₁ to C₆ substituted alkyl, C₁ to C₇ alkoxy, C₁ to C₇ substituted alkoxy, C₁ to C₇ acyl, C₁ to C₇ substituted acyl, C₁ to C₇ acyloxy, carboxy, protected carboxy, carboxymethyl, protected carboxymethyl, hydroxymethyl, protected hydroxymethyl, amino, protected amino, (monosubstituted)amino, protected (monosubstituted)amino, (disubstituted)amino, carboxamide, protected carboxamide, N-(C₁ to C₆ alkyl)carboxamide, protected N-(C₁ to C₆ alkyl)carboxamide, N, N-di(C₁ to C₆ alkyl)carboxamide, trifluoromethyl, N-((C₁ to C₆ alkyl)sulfonyl)amino, N- (phenylsulfonyl)amino or phenyl, and

 R_5 is H, C_1 to C_8 alkyl, C_1 to C_8 substituted alkyl.

4. A compound according to claim 1 wherein R₆ and R₇ are taken together with nitrogen to form the heterocycle according to the following formula (3)

$$\bigvee^{\mathsf{N}}$$

5. A compound according to any of claims 1 to 3 of the following formula (4)

formula (4)

6. A compound according to any of claims 1 to 3 of the following formula (5)

formula (5)

7. A compound according to any of claims 1 to 3 of the following formula (6)

formula (6)

8. A compound according to any of claims 1 and 4 wherein R_6 and R_7 are taken together with nitrogen to form the heterocycle according to the following formula (7)

formula (7)

9. A compound according to claim1 according to the following formula (8) formula (8)

10. A compound according to claims 1 and 4 according to the following formula (8)

- 11. A compound according to any of claims 1 to 10 wherein said compound is capable of binding the NR1H3 receptor protein or a portion thereof according to SEQ ID NO. 1 or a mammalian homologue thereof.
- 12. A compound according to any of claims 1 to 10 wherein said compound is capable of binding the NR1H2 receptor protein or a portion thereof or a mammalian homologue thereof.
- 13. Use of a compound according to any of claims 1 to 12 as a medicament
- 14. A method for prevention or treatment of a NR1H3 and/or NR1H2 receptor protein mediated disease or NR1H3 and/or NR1H2 receptor protein homologue mediated disease or condition in a mammal comprising administering a therapeutically effective amount of a compound according to any of claims 1 to 12, wherein the prevention or treatment is directly or indirectly accomplished through the binding of the compound according claims 1 to 13 to the NR1H3 and/or NR1H2 receptor proteins or to the NR1H3 and/or NR1H2 receptor protein homologues.
- 15. A method for prevention or treatment of a NR1H3 receptor protein and/or NR1H2 receptor protein mediated disease or condition according to claim 14, wherein said mammal is a human.

- 16. A method for regulating the cholesterol synthesis and/or transport in a mammal which comprises activating the NR1H3 and/or NR1H2 receptors with a therapeutically effective amount of a compound according to claims 1 to 12.
- 17. A method of treating in a mammal a disease which is affected by cholesterol, triglyceride, or bile acid levels comprising administering to a mammal in need of such treatment a therapeutically effective amount of a compound according to claims 1 to 12.
- 18. A method of treating atherosclerosis, alzheimers disease, lipid disorders, obesity or a cardiovascular disorder in a mammal, in particular a human, comprising administering to a mammal in need of such treatment a therapeutically effective amount of a compound according to claims 1 to 12.
- 19. A method according to any of claims 14 to 18 wherein the expression of ABCA1 and/or ABCG1 and/or ABCG5 and/or ABCG8 is increased.
- 20. A method according to any of claims claim 14 to 19 wherein the expression of the cholesterol 7 α hydroxylase and/or the activity of the cholesteryl ester transfer protein is increased.
- 21. A method of blocking in a mammal the cholesterol or fatty acid absorption in the intestine of a mammal in need of such blocking comprising administering to a mammal in need of such treatment a therapeutically effective amount of a compound according to claims 1 to 12.
- 22. A method for treating obesity in a mammal comprising administering a therapeutically effective amount of a compound according to any of claims 1 to 12.
- 23. A method of modulating a gene whose expression is regulated by the NR1H3 and/or NR1H2 receptor in a mammal comprising administering a therapeutically effective amount of a compound according to claims 1 to 10.

- 24. A method according to any of claims claim 14 to 19 wherein the expression of the cholesterol 7 α hydroxylase and/or the activity of the cholesteryl ester transfer protein is enhanced.
- 25. Use of a compound according to any of claims 1 to 12 wherein the mammal is a human
- 26. Use of a compound according to any of claims 1 to 12 for the manufacture of a medicament for the prevention or treatment of a NR1H3 and/or NR1H2 receptor protein or NR1H3 and/or NR1H2 receptor protein homologue mediated disease or condition in a mammal wherein the prevention or treatment is directly or indirectly accomplished through the binding of the compound according claims 1 to 8 to the NR1H3 and/or NR1H2 receptor protein or NR1H3 and/or NR1H2 receptor protein homologue.
- 27. Use of a compound according to any of claims 1 to 12 for the manufacture of a medicament for prevention or treatment of a NR1H3 and/or NR1H2 receptor protein mediated disease or condition according to claim 26, wherein the mammal is a human.
- 28. Use of a compound according to any of claims 1 to 12 for the manufacture of a medicament for regulating the cholesterol transport system in a mammal by activating the NR1H3 and/or NR1H2 receptor.
- 29. Use of a compound according to any of claims 1 to 12 for the manufacture of a medicament for regulating levels of cholesterol, triglyceride, and/or bile acid.
- 30. Use of a compound according to any of claims 1 to 12 for the manufacture of a medicament for treating in a mammal atherosclerosis, alzheimer disease, gallstone disease, lipid disorders, obesity or a cardiovascular disorder.
- 31. Use of a compound according to any of claims 1 to 12 for the manufacture of a medicament capable for blocking in a mammal the cholesterol and/or fatty acid absorption in the intestine.

- 32. Use of the compound according to any of claims 1 to 12 for the manufacture of a medicament for treating obesity in a mammal.
- 33. Use of a compound according to any of claims 1 to 12 for the manufacture of a medicament for modulating a gene whose expression is regulated by the NR1H3 and/or NR1H2 receptor.
- 34. Use of a compound according to any of claims 1 to 12 in a mammal for the selective up-regulation of one or more genes selected from the group consisting of ABCA1, ABCG1, ABCG5 and ABCG8 and a down-regulation of one or more of the genes selected from the group comprising FAS and SREBP-1c, said compound showing a larger difference in regulation of the two groups of genes when compared with the regulatory behavior of T0901317 on both groups of genes.
- 35. Use of a compound according to claims 28, 30, 31, 32, and 34 wherein the mammal is a human.

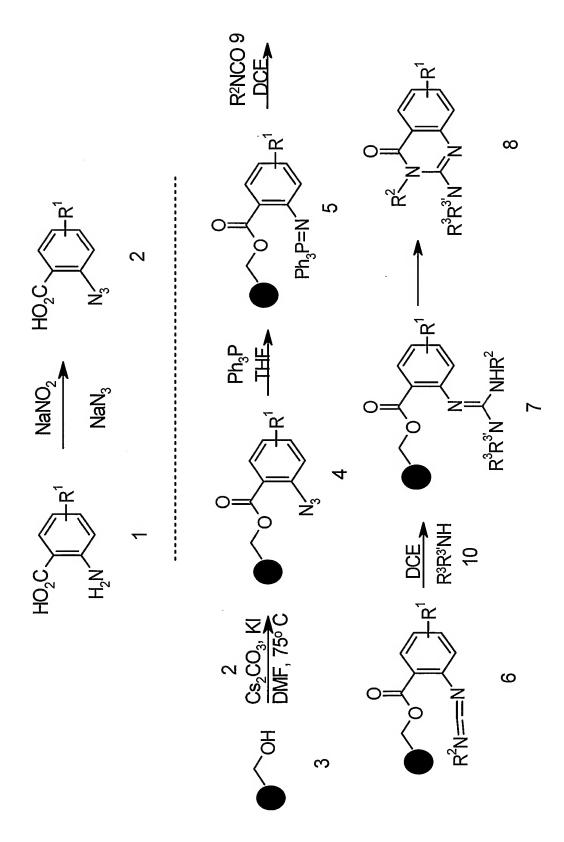


Fig.1

Fig. 2 : Measurement parameters employed by the Wallace VICTOR2V $^{\text{TM}}$ Multilabel Counter:

Counter:
Number of repeats 1
plate: GREINER FIA-Plate black 384 well med. binding
Measurement height 3.50 mm
Label technology TR-F Lance
Emission filter name D615
Emission filter slot
Emission aperture Normal
Excitation filter D340
Delay 50 μs
Window time 400 μs
Cycle 1000 μs
Light integrator capacitors 1
Light integrator ref. level 95
Flash energy area High
Flash energy level 223
Flash absorbance measurement No
Beam Normal
Label technology TR-F Lance
Emission filter name D665
Emission filter slot A8
Emission aperture Normal
Excitation filter D340
Delay 50 μs
Window time 400 μs
Cycle 1000 μs
Light integrator capacitors 1
Light integrator ref. level 95
Flash energy area High
Flash energy level 223
Flash absorbance measurement No
Beam Normal

3/16

Fig. 3 A and B:

A

```
1 mslwlgapvp dippdsavel wkpgaqdass qaqggsscil reearmphsa ggtagvglea 61 aeptalltra eppsepteir pqkrkkgpap kmlgnelcsv cgdkasgfhy nvlscegckg 121 ffrrsvikga hyichsgghc pmdtymrrkc qecrlrkcrq agmreecvls eeqirlkklk 181 rqeeeqahat slpprrsspp qilpqlspeq lgmieklvaa qqqcnrrsfs drlrvtpwpm 241 apdphsrear qqrfahftel aivsvqeivd fakqlpgflq lsredqiall ktsaievmll 301 etsrrynpgs esitflkdfs ynredfakag lqvefinpif efsramnelq lndaefalli 361 aisifsadrp nvqdqlqver lqhtyvealh ayvsihhphd rlmfprmlmk lvslrtlssv 421 hseqvfalrl qdkklpplls eiwdvhe
```

В

```
1 atgtccttgt ggctgggggc ccctgtgcct gacattcctc ctgactctgc ggtggagctg 61 tggaagccag gcgcacagga tgcaagcagc caggcccagg gaggcagcag ctgcatcctc
 121 agagaggaag ccaggatgcc ccactctgct gggggtactg caggggtggg gctggaggct
 181 gcagagccca cagccctgct caccagggca gagccccctt cagaacccac agagatccgt
 241 ccacaaaagc ggaaaaaggg gccagcccc aaaatgctgg ggaacgagct atgcagcgtg
 301 tgtggggaca aggcctcggg cttccactac aatgttctga gctgcgaggg ctgcaaggga 361 ttcttccgcc gcagcgtcat caagggageg cactacatct gccacagtgg cggccactgc
 421 cccatggaca cctacatgcg tcgcaagtgc caggagtgtc ggcttcgcaa atgccgtcag
 481 gctggcatgc gggaggagtg tgtcctgtca gaagaacaga tccgcctgaa gaaactgaag
541 cggcaagagg aggaacaggc tcatgccaca tccttgcccc ccaggcgttc ctcacccccc
 601 caaatcctgc cccagctcag cccggaacaa ctgggcatga tcgagaagct cgtcgctgcc
 661 cagcaacagt gtaaccggcg ctccttttct gaccggcttc gagtcacgcc ttggcccatg
 721 gcaccagate eccatageeg ggaggeegt eageageget ttgeecaett caetgagetg
 781 gccatcgtct ctgtgcagga gatagttgac tttgctaaac agctacccgg cttcctgcag
841 ctcagccggg aggaccagat tgccctgctg aagacctctg cgatcgaggt gatgcttctg
 901 gagacatete ggaggtacaa eeetgggagt gagagtatea eetteeteaa ggattteagt
 961 tataaccggg aagactttgc caaagcaggg ctgcaagtgg aattcatcaa ccccatcttc
1021 gagttctcca gggccatgaa tgagctgcaa ctcaatgatg ccgagtttgc cttgctcatt
1081 gctatcagca tcttctctgc agaccggccc aacgtgcagg accagctcca ggtggagagg
1141 ctgcagcaca catatgtgga agccctgcat gcctacgtct ccatccacca tccccatgac
1201 cgactgatgt tcccacggat gctaatgaaa ctggtgagcc tccggaccct gagcagcgtc
1261 cactcagage aagtgtttgc actgcgtctg caggacaaaa agctcccace gctgctctct
1321 gagatctggg atgtgcacga atga
```

Fig. 3 C and D:

\mathbf{C}

PVLPSSESFT TRQDLQGKIT SLDTSTMRAA MKPGWEDLVR RCIQKFHAQH EGESVSYAKR 120 HHHEVLRQGL AFSQ1YRFSL SDGTLVAAQT KSKLIRSQTT NEPQLVISLH MLHREQNVCV 180 MNPDLTGQTM GKPLNPISSN SPAHQALCSG NPGQDMTLSS NINFFINGPK EQMGMPMGRF 240 GGSGGMNHVS GMQATTPQGS NYALKMNSPS QSSPGMNPGQ PTSMLSPRHR MSPGVAGSPR 300 LPPSQFSPAG SLHSPVGVCS STGNSHSYTN SSLNALQALS EGHGVSLGSS LASPDLKMGN 360 LQNSPVNMNP PPLSKMGSLD SKDCFGLYGE PSEGTTGQAE SSCHPGEQKE TNDPNLPPAV 420 SSERADGQSR LHDSKGQTKL LQLLTTKSDQ MEPSPLASSL SDTNKDSTGS LPGSGSTHGT 480 SLKEKHKILH RLLQDSSSPV DLAKLTAEAT GKDLSQESSS TAPGSEVTIK QEPVSPKKKE 540 NALLRYLLDK DDTKDIGLPE ITPKLERLDS KTDPASNTKL IAMKTEKEEM SFEPGDQPGS 600 ELDNLEEILD DLQNSQLPQL FPDTRPGAPA GSVDKQAILN DLMQLTAENS VPTPVGAQKT 660 ALRISQSTFN NPRPGQLGRL LPNQNLPLDI TLQSPTGAGP FPPIRNSSPY SVIPQPGMMG 720 NQCMIGNQGN LGNSSTGMIG NSASRPTMPS GEWAPQSSAV RVTCAATTSA MNRPVQGGMI 780 NQCMIGNQGN LGNSSTGMIG NSASRPTMPS GEWAPQSSAV RVTCAATTSA MNRPVQGGMI 780 RNPAASIPMR PSSQPGQRQT LQSQVMNIGP SELEMNMGGP QYSQQQAPPN QTAPWPESIL 840 PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFPQQYASQA QMAQGSYSPM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMNQIS NVSNVNLTLR 1020 PGVPTQAPIN AQMLAQRQRE ILNQHLRQRQ MHQQQVQVQR TLMMRGGGLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFPF PNYGISQQF DPGFTGATTP QSPLMSPRMA HTQSPMMQPS 1140 QANPAYQAPS DINGWAQGMM GGNSMFSQQS PPHFGQQANT SMYSNNMIN VSMATNTGGM 1260 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	MLVKPLPDSE	EEGHDNQEAH	QKYETMQCFA	VSQPKSIKEE	GEDLQSCLIC	VARRVPMKER	60
MNPDLTGQTM GKPLNPISSN SPAHQALCSG NPGQDMTLSS NINFPINGPK EQMGMPMGRF GSGGGMNHVS GMQATTPQGS NYALKMNSPS QSSPGMNPGQ PTSMLSPRHR MSPGVAGSPR 300 IPPSQFSPAG SLHSPVGVCS STGNSHSYTN SSINALQALS EGHGVSLGSS LASPDLKMGN 360 LQNSPVNMNP PPLSKMGSLD SKDCFGLYGE PSEGTTGQAE SCCHPGEQKE TNDPNLPPAV 420 SSERADGQSR LHDSKGQTKL LQLLTTKSDQ MEPSPLASSL SDTNKDSTGS LPGSGSTHGT 480 SLKEKKKILH RLLQDSSSPV DLAKLTAEAT GKDLSQESSS TAPGSEVTK QEPVSPKKKE 540 NALLRYLLDK DDTKDIGLPE ITPKLERLDS KTDPASNTKL IAMKTEKEEM SFEPGDQPGS 600 ELDNLEEILD DLQNSQLPQL FPDTRPGAPA GSVDKQAIIN DLMQLTAENS PVTPVGAQKT 660 ALRISQSTFN NPRPGQLGRL LPNQNLPLDI TLQSPTGAGP FPPIRNSSPY SVIPQPGMMG 720 RQGMIGNQGN LGNSSTGMIG NSASRPTMPS GEWAPQSSAV RVTCAATTSA MNRPVQGGMI 780 RNPAASIPMR PSSQPGQRQT LQSQVMNIGP SELEMNMGGP QYSQQQAPPN QTAPWPESIL 840 PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFPQQYASQA QMAQGSYSPM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRFTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMNQIS NVSNVNLTLR 1020 PGVPTQAPIN AQMLAQRQFF PNYGISQQP DPGFTGATTP SPLMSRPAM HTQSPMMQPS 1080 MPATMSNPRI PQANAQQFPF PNYGISQQP DPGFTGATTP SPLMSRPRM HTQSPMMQPS 1140 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	PVLPSSESFT	TRQDLQGKIT	SLDTSTMRAA	MKPGWEDLVR	RCIQKFHAQH	EGESVSYAKR	120
GGSGGMNHVS GMQATTPQGS NYALKMNSPS QSSPGMNPGQ PTSMLSPRHR MSPGVAGSPR 300 IPPSQFSPAG SLHSPVGVCS STGNSHSYTN SSLNALQALS EGHGVSLGSS LASPDLKMGN 360 LQNSPVNMNP PPLSKMGSLD SKDCFGLYGE PSEGTTGQAE SSCHPGEQKE TNDPNLPPAV 420 SSERADGQSR LHDSKGQTKL LQLLTTKSDQ MERSPLASSL SDTNKDSTGS LPGSGSTHGT 480 SLKEKHKILH RLLQDSSSPV DLAKLTAEAT KØLDSQSSS TAPGSEVTIK QEPVSPKKKE 540 NALLRYLLDK DDTKDIGLPE ITPKLERLDS KTDPASNTKL IAMKTEKEEM SFEPGDQPGS 600 ELDNLEEILD DLQNSQLPQL FPDTRPGAPA GSVDKQAIIN DLMQLTAENS PVTPVGAQKT 660 ALRISQSTFN NPRPGQLGRL LPNQNLPLDI TLQSPTGAGP FPPIRNSSPY SVIPQPGMMG 720 NQGMIGNQGN LGNSSTGMIG NSASRPTMPS GEWAPQSSAV RVTCAATTSA MNRPVQGGMI 780 RNPAASIPMR PSSQPGQRQT LQSQVMNIGP SELEMNMGGP QYSQQQAPPN QTAPWPESIL 840 PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFPQQYASQA QMAQGSYSPM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMNQIS NVSNVNLTLR 1020 PGVPTQAPIN AQMLAQRQRE ILNQHLRQRQ MHQQQVQQR TLMMRGGGLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFFF PPNYGISQQP DPGFTGATTP QSPLMSPRMA HTQSPMMQOS 1140 QANPAYQAPS DINGWAQGNM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	HHHEVLRQGL	AFSQIYRFSL	SDGTLVAAQT	KSKLIRSQTT	NEPQLVISLH	MLHREQNVCV	180
IPPSQFSPAG SLHSPVGVCS STGNSHSYTN SSLNALQALS EGHGVSLGSS LASPDLKMGN 360 LQNSPVNMNP PPLSKMGSLD SKDCFGLYGE PSEGTTGQAE SSCHPGEQKE TNDPNLPPAV 420 SSERADGQSR LHDSKGQTKL LQLLTTKSDQ MEPSPLASSL SDTNKDSTGS LPGSGSTHGT 480 SLKEKHKILH RLLQDSSSPV DLAKLTAEAT GKDLSQESSS TAPGSEVTIK QEPVSPKKKE 540 NALLRYLLDK DDTKDIGLPE ITPKLERLDS KTDPASNTKL IAMKTEKEEM SFEPGDQPGS 600 ELDNLEEILD DLQNSQLPQL FPDTRPGAPA GSVDKQAIIN DLMQLTAENS PVTPVGAQKT 660 ALRISQSTFN NPRPGQLGRL LPNQNLPLDI TLQSPTGAGP FPPIRNSSPY SVIPQPGMMG 720 NQGMIGNQGN LGNSSTGMIG NSASRPTMPS GEWAPQSSAV RVTCAATTSA MNRPVQGGMI 780 RNPAASIPMR PSSQPGQRQT LQSQVMNIGP SELEMNMGGP QYSQQQAPPN QTAPWPESIL 840 PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFFQQYASQA QMAQGSYSPM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMNQIS NVSNVNLTLR 1020 PGVPTQAPIN AQMLAQRQRE ILNQHLRQRQ MHQQQVQQR TLMMRGGGLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFFF PPNYGISQQP DPGFTGATTP QSPLMSPRMA HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGNM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	MNPDLTGQTM	GKPLNPISSN	SPAHQALCSG	NPGQDMTLSS	NINFPINGPK	EQMGMPMGRF	240
LQNSPVNMNP PPLSKMGSLD SKDCFGLYGE PSEGTTGQAE SSCHPGEQEE TNDPNLPPAV 420 SSERADGQSR LHDSKGQTKL LQLLTTKSDQ MEPSPLASSL SDTNKDSTGS LPGSGSTHGT 480 SLKEKHKILH RLLQDSSSPV DLAKLTAEAT GKDLSQESSS TAPGSEVTIK QEPVSPKKKE 540 NALLRYLLDK DDTKDIGLPE ITPKLERLDS KTDPASNTKL IAMKTEKEEM SFEPGDQPGS 600 ELDNLEEILD DLQNSQLPQL FPDTRPGAPA GSVDKQAIIN DLMQLTAENS PVTPVGAQKT 660 ALRISQSTFN NPRPGQLGRL LPNQNLPLDI TLQSPTGAGP FPPIRNSSPY SVIPQPGMMG 720 NQCMIGNQGN LGNSSTGMTG NSASRPTMPS GEWAPQSSAV RVTCAATTSA MNRPVQGGMI 780 RNPAASIPMR PSSQPGQRQT LQSQVMNIGP SELEMNMGGP QYSQQQAPPN QTAPWPESIL 840 PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFFQQYASQA QMAQGSYSPM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAGQ NRQPLMQLS NVSNVNLTLR 1020 PGVPTQAPIN AQMLAQRQRE ILNQHLRQRQ MHQQQVQQR TLMMRGGGLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFPF PPNYGISQQP DPGFTGATTP QSPLMSPRMA HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGMM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	GGSGGMNHVS	GMQATTPQGS	NYALKMNSPS	QSSPGMNPGQ	PTSMLSPRHR	MSPGVAGSPR	300
SSERADGQSR LHDSKGQTKL LQLLTTKSDQ MEPSPLASSL SDTNKDSTGS LPGSGSTHGT 480 SLKEKHKILH RLLQDSSSPV DLAKLTAEAT GKDLSQESSS TAPGSEVTIK QEPVSPKKKE 540 NALLRYLLDK DDTKDIGLPE ITPKLERLDS KTDPASNTKL IAMKTEKEEM SFEPGDQPGS 600 ELDNLEEILD DLQNSQLPQL FPDTRPGAPA GSVDKQAIIN DLMQLTAENS PVTPVGAQKT 660 ALRISQSTFN NPRPGQLGRL LPNQNLPLDI TLQSPTGAGP FPPIRNSSPY SVIPQPGMMG 720 NQCMIGNQGN LGNSSTGMIG NSASRPTMPS GEWAPQSSAV RVTCAATTSA MNRPVQGGMI 780 RNPAASIPMR PSSQPGQRQT LQSQVMNIGP SELEMNMGGP QYSQQQAPPN QTAPWPESIL 840 PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFFQQYASQA QMAQGSYSPM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMQIS NVSNVNLTLR 1020 PGVPTQAPIN AQMLAQRQRE ILNQHLRQRQ MHQQQVQQR TLMMRGGGLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFPF PPNYGISQQP DPGFTGATTP QSPLMSPRMA HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGMM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	IPPSQFSPAG	SLHSPVGVCS	STGNSHSYTN	SSLNALQALS	EGHGVSLGSS	LASPDLKMGN	360
SLKEKHKILH RLLQDSSSPV DLAKLTAEAT GKDLSQESSS TAPGSEVTIK QEPVSPKKKE 540 NALLRYLLDK DDTKDIGLPE TTPKLERLDS KTDPASNTKL IAMKTEKEEM SFEPGDQPGS 600 ELDNLEEILD DLQNSQLPQL FPDTRPGAPA GSVDKQAIIN DLMQLTAENS PVTPVGAQKT 660 ALRISQSTFN NPRPGQLGRL LPNQNLPLDI TLQSPTGAGP FPPIRNSSPY SVIPQPGMMG 720 NQGMIGNQGN LGNSSTGMIG NSASRPTMPS GEWAPQSSAV RVTCAATTSA MNRPVQGGMI 780 RNPAASIPMR PSSQPGQRQT LQSQVMNIGP SELEMNMGGP QYSQQQAPPN QTAPWPESIL 840 PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFPQQYASQA QMAQGSYSPM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMNQIS NVSNVNLTLR 1020 PGVPTQAPIN AQMLAQRQRE ILNQHLRQRQ MMQQQVQQR TLMMRGGQLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFPF PPNYGISQQP DPGFTGATTP SPLMSRPRM HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGMM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	LQNSPVNMNP	PPLSKMGSLD	SKDCFGLYGE	PSEGTTGQAE	SSCHPGEQKE	TNDPNLPPAV	420
NALLRYLLDK DDTKDIGLPE ITPKLERLDS KTDPASNTKL IAMKTEKEEM SFEPGDQPGS 600 ELDNLEEILD DLQNSQLPQL FPDTRPGAPA GSVDKQAIIN DLMQLTAENS PVTPVGAQKT 660 ALRISQSTFN NPRPGQLGRL LPNQNLPLDI TLQSPTGAGG FPPIRNSSFY SVIPQPGMMG 720 NQGMIGNQGN LGNSSTGMIG NSASRPTMPS GEWAPQSSAV RVTCAATTSA MNRPVQGGMI 780 RNFAASIPMR PSSQPGQRQT LQSQVMNIGP SELEMNMGGP QVSQQQAPPN QTAPWPESIL 840 PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFPQQYASQA QMAQGSYSPM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMNQIS NVSNVNLTLR 1020 PGVPTQAPIN AQMLAQRQRE ILNQHLRQRQ DLQHRLQAQQ NRQPLMQIS NVSNVNLTLR 1020 PGVPTQAPIN PQANAQQFPF PPNYGISQQP DPGFTGATTP QSPLMSPRMA HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGMM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	SSERADGQSR	LHDSKGQTKL	LQLLTTKSDQ	${\tt MEPSPLASSL}$	SDTNKDSTGS	LPGSGSTHGT	480
ELDNLEEILD DLQNSQLPQL FPDTRPGAPA GSVDKQAIIN DLMQLTAENS PVTPVGAQKT 660 ALRISQSTFN NPRPGQLGRL LPNQNLPLDI TLQSPTGAGP FPPIRNSSPY SVIPQPGMMG 720 NQGMIGNQGN LGNSSTGMIG NSASRPTMPS GEWAPQSSAV RVTCAATTSA MNRPVQGGMI 780 RNPAASIPMR PSSQPGQRQT LQSQVMNIGP SELEMNMGGP QYSQQQAPPN QTAPWPESIL 840 PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFPQQYASQA QMSQGSYSPM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMNQIS NVSNVNLTLR 1020 RPSYATLRMQ PAGNLAQRQRE ILNQHLRQRQ MHQQQQVQQR TLMMRGGGLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFFF PPNYGISQQP DPGFTGATTP QSPLMSPRMA HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGNM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	SLKEKHKILH	RLLQDSSSPV	DLAKLTAEAT	GKDLSQESSS	TAPGSEVTIK	QEPVSPKKKE	540
ALRISQSTFN NPRPGQLGRL LPNQNLPLDI TLQSPTGAGP FPPIRNSSPY SVIPQPGMMG 720 NQGMIGNQGN LGNSSTGMIG NSASRPTMPS GEWAPQSSAV RVTCAATTSA MNRPVQGGMI 780 RNPAASIPMR PSSQPGQRQT LQSQVMNIGP SELEMNMGGP QYSQQQAPPN QTAPWPESIL 840 PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFPQQYASQA QMSQGSYSPM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMNQIS NVSNVNLTLR 1020 RPSYATLRMQ PRGLRPTGL LNQHLRQRQ MHQQQQVQQR TLMMRGGGLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFFF PPNYGISQQP DFGFTGATTP QSPLMSFRMA HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGNM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	NALLRYLLDK	DDTKDIGLPE	ITPKLERLDS	KTDPASNTKL	IAMKTEKEEM	SFEPGDQPGS	600
NQGMIGNQGN LGNSSTGMIG NSASRPTMPS GEWAPQSSAV RVTCAATTSA MNRPVQGGMI 780 RNPAASIPMR PSSQPGQRQT LQSQVMNIGP SELEMNMGGP QYSQQQAPPN QTAPWPESIL 840 PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFFQQYASQA QMAQGSYSFM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMNQIS NVSNVNLTLR 1020 PGVPTQAPIN AQMLAQRQRE ILNQHLRQRQ MHQQQVQQR TLMMRGGGLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFFF PPNYGISQQP DFGFTGATTP QSPLMSFRMA HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGNM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	ELDNLEEILD	DLQNSQLPQL	FPDTRPGAPA	GSVDKQAIIN	DLMQLTAENS	PVTPVGAQKT	660
RNPAASIPMR PSSQPGQRQT LQSQVMNIGP SELEMNMGGP QYSQQQAPPN QTAPWPESIL 840 PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFFQQYASQA QMAQGSYSPM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMNQIS NVSNVNLTLR 1020 PGVPTQAPIN AQMLAQRQRE ILNQHLRQRQ MHQQQQVQQR TLMMRGGGLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFPF PPNYGISQQP DPGFTGATTP QSPLMSPRMA HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGMM GGNSMFSQQS PPHFGQQANT SMYSNNMIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	ALRISQSTFN	NPRPGQLGRL	LPNQNLPLDI	TLQSPTGAGP	FPPIRNSSPY	SVIPQPGMMG	720
PIDQASFASQ NRQPFGSSPD DLLCPHPAAE SPSDEGALLD QLYLALRNFD GLEEIDRALG 900 IPELVSQSQA VDPEQFSSQD SNIMLEQKAP VFPQQYASQA QMAQGSYSPM QDPNFHTMGQ 960 RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMNQIS NVSNVNLTLR 1020 PGVPTQAPIN AQMLAQRQRE ILNQHLRQRQ MHQQQQVQR TLMMRGGQLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFPF PPNYGISQQP DPGFTGATTP QSPLMSPRMA HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGMM GGNSMFSQQS PPHFGQQANT SMYSNNMIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	NQGMIGNQGN	LGNSSTGMIG	NSASRPTMPS	GEWAPQSSAV	RVTCAATTSA	MNRPVQGGMI	780
IPELVSQSQAVDPEQFSSQDSNIMLEQKAPVFPQQYASQAQMAQGSYSPMQDPNFHTMGQ960RPSYATLRMQPRPGLRPTGLVQNQPNQLRLQLQHRLQAQQNRQPLMNQISNVSNVNLTLR1020PGVPTQAPINAQMLAQRQREILNQHLRQRQMHQQQQVQQRTLMMRGQGLNMTPSMVAPSG1080MPATMSNPRIPQANAQQFPFPPNYGISQQPDPGFTGATTPQSPLMSPRMAHTQSPMMQQS1140QANPAYQAPSDINGWAQGNMGGNSMFSQQSPPHFGQQANTSMYSNNMNINVSMATNTGGM1200SSMNQMTGQISMTSVTSVPTSGLSSMGPEQVNDPALRGGNLFPNQLPGMDMIKQEGDTTR1260	RNPAASIPMR	PSSQPGQRQT	LQSQVMNIGP	SELEMNMGGP	QYSQQQAPPN	QTAPWPESIL	840
RPSYATLRMQ PRPGLRPTGL VQNQPNQLRL QLQHRLQAQQ NRQPLMNQIS NVSNVNLTLR 1020 PGVPTQAPIN AQMLAQRQRE ILNQHLRQRQ MHQQQQVQQR TLMMRGQGLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFPF PPNYGISQQP DPGFTGATTP QSPLMSPRMA HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGNM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	PIDQASFASQ	NRQPFGSSPD	DLLCPHPAAE	SPSDEGALLD	QLYLALRNFD	GLEEIDRALG	900
PGVPTQAPIN AQMLAQRQRE ILNQHLRQRQ MHQQQQVQQR TLMMRGQGLN MTPSMVAPSG 1080 MPATMSNPRI PQANAQQFPF PPNYGISQQP DPGFTGATTP QSPLMSPRMA HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGNM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	IPELVSQSQA	VDPEQFSSQD	SNIMLEQKAP	VFPQQYASQA	QMAQGSYSPM	QDPNFHTMGQ	960
MPATMSNPRI PQANAQQFPF PPNYGISQQP DPGFTGATTP QSPLMSPRNA HTQSPMMQQS 1140 QANPAYQAPS DINGWAQGNM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	RPSYATLRMQ	PRPGLRPTGL	VQNQPNQLRL	QLQHRLQAQQ	NRQPLMNQIS	NVSNVNLTLR	1020
QANPAYQAPS DINGWAQGNM GGNSMFSQQS PPHFGQQANT SMYSNNMNIN VSMATNTGGM 1200 SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	PGVPTQAPIN	AQMLAQRQRE	ILNQHLRQRQ	MHQQQQVQQR	TLMMRGQGLN	MTPSMVAPSG	1080
SSMNQMTGQI SMTSVTSVPT SGLSSMGPEQ VNDPALRGGN LFPNQLPGMD MIKQEGDTTR 1260	MPATMSNPRI	PQANAQQFPF	PPNYGISQQP	DPGFTGATTP	QSPLMSPRMA	HTQSPMMQQS	1140
	QANPAYQAPS	DINGWAQGNM	GGNSMFSQQS	PPHFGQQANT	SMYSNNMNIN	VSMATNTGGM	1200
KYC 1263	SSMNQMTGQI	SMTSVTSVPT	SGLSSMGPEQ	VNDPALRGGN	LFPNQLPGMD	MIKQEGDTTR	1260
	KYC						1263

D

1 ggcggccgca gcctcggcta cagcttcggc ggcgaaggtc agcgccgacg gcagccggca 61 cctgacggcg tgaccgaccc gagccgattt ctcttggatt tggctacaca cttatagatc 121 ttctgcactg tttacaggca cagttgctga tatgtgttca agatgagtgg gatgggagaa 181 aatacctctg acccctccag ggcagagaca agaaagcgca aggaatgtcc tgaccaactt 241 ggacccagcc ccaaaaggaa cactgaaaaa cgtaatcgtg aacaggaaaa taaatatata 301 gaagaacttg cagagttgat ttttgcaaat tttaatgata tagacaactt taacttcaaa 361 cctgacaaat gtgcaatctt aaaagaaact gtgaagcaaa ttcgtcagat caaagaacaa 421 gagaaagcag cagctgccaa catagatgaa gtgcagaagt cagatgtatc ctctacaggg 481 cagggtgtca tcgacaagga tgcgctqggg cctatgatgc ttgaqgccct tqatgggttc 541 ttctttgtag tgaacctgga aggcaacgtt gtgtttgtgt cagagaatgt gacacagtat 601 ctaaggtata accaagaaga gctgatgaac aaaagtgtat atagcatcit gcatgttggg 661 gaccacacgg aatttgtcaa aaacctgctg ccaaagtcta taggtaaatg ggggatcttg 721 gtctggcgaa cctccgaggc ggaacagcca taccttcaat tgtcggatgc tggtaaaacc 781 tttacctgat tcagaagagg agggtcatga taaccaggaa gctcatcaga aatatgaaac 841 tatgcagtgc ttcgctgtct ctcaaccaaa gtccatcaaa gaagaaggag aagatttgca 901 gtcctgcttg atttgcgtgg caagaagagt tcccatgaag gaaagaccag ttcttcctc 961 atcagaaagt tttactactc gccaggatct ccaaggcaag atcacgtctc tggataccag 1021 caccatgaga gcagccatga aaccaggctg ggaggacctg gtaagaaggt gtattcagaa 1081 gttccatgcg cagcatgaag gagaatctgt gtcctatgct aagaggcatc atcatgaagt 1141 actgagacaa ggattggcat tcagtcaaat ctatcgtttt tccttgtctg atggcactct 1201 tgttgctgca caaacgaaga gcaaactcat ccgttctcag actactaatg aacctcaact 1261 tgtaatatct ttacatatgc ttcacagaga gcagaatgtg tgtgtgatga atccggatct 1321 gactggacaa acgatgggga agccactgaa tccaattagc tctaacagcc ctgcccatca 1381 ggccctgtgc agtgggaacc caggtcagga catgaccctc agtagcaata taaattttcc 1441 cataaatggc ccaaaggaac aaatgggcat gcccatgggc aggtttggtg gttctggggg 1501 aatgaaccat gtgtcaggca tgcaagcaac cactcctcag ggtagtaact atgcactcaa 1561 aatgaacago coctoacaaa goagoootgg catgaatoca ggacagooca cotocatgot 1621 ttcaccaagg catcgcatga gccctggagt ggctggcagc cctcgaatcc cacccagtca 1681 gttttcccct gcaggaagct tgcattcccc tgtgggagtt tgcagcagca caggaaatag 1741 ccatagttat accaacagct ccctcaatgc acttcaggcc ctcagcgagg ggcacggggt 1801 ctcattaggg tcatcgttgg cttcaccaga cctaaaaatg ggcaatttgc aaaactcccc 1861 agttaatatg aatcetecee cacteageaa gatgggaage ttggacteaa aagactgttt 1921 tggactatat ggggagcct ctqaaggtac aactggacaa gcagagagca gctgccatcc 1981 tggagagcaa aaggaaacaa atgaccccaa cctgcccccg gccgtgagca gtgagagagc 2041 tgacgggcag agcagactgc atgacagcaa agggcagacc aaactcctgc agctgctgac 2101 caccaaatct gatcagatgg agccctcgcc cttagccagc tctttgtcgg atacaaacaa 2161 agactecaca ggtagettge etggttetgg gtetacacat ggaacetege teaaggagaa 2221 qcataaaatt ttqcacagac tcttqcaqga cagcagttcc cctgtggact tggccaagtt 2281 aacagcagaa gccacaggca aagacctgag ccaggagtcc agcagcacag ctcctggatc 2341 agaagtgact attaaacaag agccggtgag ccccaagaag aaagagaatg cactacttcg 2401 ctatttgcta gataaagatg atactaaaga tattggttta ccagaaataa cccccaaact 2461 tgagagactg gacagtaaga cagatcctgc cagtaacaca aaattaatag caatgaaaac 2521 tgagaaggag gagatgagct ttgagcctgg tgaccagcct ggcagtgagc tggacaactt 2581 ggaggagatt ttggatgatt tgcagaatag tcaattacca cagcttttcc cagacacgag 2641 gccaggcgcc cctgctggat cagttgacaa gcaagccatc atcaatgacc tcatgcaact 2701 cacagetgaa aacageeetg teacacetgt tggageeeag aaaacageae tgegaattte

Fig. 3 D (continued):

2761	acagagcact	tttaataacc	cacgaccagg	gcaactgggc	aggttattqc	caaaccagaa
2821	tttaccactt	gacatcacat	tgcaaagccc	aactggtgct	ggacctttcc	caccaatcag
2881	aaacagtagt	ccctactcag	tgatacctca	gccaggaatg	atgggtaatc	aagggatgat
2941	aggaaaccaa	ggaaatttag	ggaacagtag	cacaggaatg	attggtaaca	gtgcttctcg
3001	gcctactatg	ccatctggag	aatgggcacc	gcagagttcg	gctgtgagag	tcacctgtgc
3061	tgctaccacc	agtgccatga	accggccagt	ccaaggaggt	atgattcgga	acccagcagc
	cagcatcccc					
	catgaatata					
3241	acaagctcct	ccaaatcaga	ctgccccatg	gcctgaaagc	atcctgccta	tagaccaggc
3301	gtcttttgcc	agccaaaaca	ggcagccatt	tggcagttct	ccagatgact	tgctatgtcc
3361	acatcctgca	gctgagtctc	cgagtgatga	gggagctctc	ctggaccagc	tgtatctggc
3421	cttgcggaat	tttgatggcc	tggaggagat	tgatagagcc	ttaggaatac	ccgaactggt
	cagccagagc					
	ggagcagaag					
3601	tagctattct	cccatgcaag	atccaaactt	tcacaccatg	ggacagcggc	ctagttatgc
3661	cacactccgt	atgcagccca	gaccgggcct	caggcccacg	ggcctagtgc	agaaccagcc
3721	aaatcaacta	agacttcaac	ttcagcatcg	cctccaagca	cagcagaatc	gccagccact
3781	tatgaatcaa	atcagcaatg	tttccaatgt	gaacttgact	ctgaggcctg	gagtaccaac
3841	acaggcacct	attaatgcac	agatgctggc	ccagagacag	agggaaatcc	tgaaccagca
3901	tcttcgacag	agacaaatgc	atcagcaaca	gcaagttcag	caacgaactt	tgatgatgag
3961	aggacaaggg	ttgaatatga	caccaagcat	ggtggctcct	agtggtatgc	cagcaactat
4021	gagcaaccct	cggattcccc	aggcaaatgc	acagcagttt	ccatttcctc	caaactacgg
4081	aataagtcag	caacctgatc	caggctttac	tggggctacg	actccccaga	gcccacttat
4141	gtcaccccga	atggcacata	cacagagtcc	catgatgcaa	cagtctcagg	ccaacccagc
4201	ctatcaggcc	ccctccgaca	taaatggatg	ggcgcagggg	aacatgggcg	gaaacagcat
4261	gttttcccag	cagtccccac	cacactttgg	gcagcaagca	aacaccagca	tgtacagtaa
4321	caacatgaac	atcaatgtgt	ccatggcgac	caacacaggt	ggcatgagca	gcatgaacca
4381	gatgacagga	cagatcagca	tgacctcagt	gacctccgtg	cctacgtcag	ggctgtcctc
	catgggtccc					
	gctgcctgga					
	ctgctgaagc					
	tggagagctg					
	gacaggcctg					
	tcttctcttg					
4801	accttagtgc	aacttagatc	tctcctgcaa	aagtaaatgt	tgacaggcaa	atttcatacc
4861	catgtcagat	tgaatgtatt	taaatgtatg	tatttaagga	gaaccatgct	cttgttctgt
4921	tcctgttcgg	ttccagacac	tggtttcttg	ctttgttttc	cctggctaac	agtctagtgc
4981	aaaagattaa	gattttatct	gggggaaaga	aaagaatttt	ttaaaaaatt	aaactaaaga
	tgttttaagc					
	tatttacaga					
	ctctaaagac					
	gagggctgtt					
	gttcacctga					
	gtttgatcag					
	ccctctctca					
	cagaaaccag					
5521	agtattgaat	ttcagaaaaa	aaaaatgcat	ttactggcaa	ggagaagagc	aaagttaagg
	cttgatacca					
	aactctggcc					
	gacagattta					
	gaagttttgt					
	gaacttatgt					
	aaaatgaggt					
	ttgtaaagga					
	ctaaaccttg					
	taggaaccaa				ccaatgaacc	ıcagaaagac
0171	tacatgttac	Laagcaggcc	acttttatgg	ccgttttt		

Fig. 3 E:

1 MSSPTTSSLD TPLPGNGPPQ PGAPSSSPTV KEEGPEPWPG GPDPDVPGTD EASSACSTDW
61 VIPDPEEPE RKRKKGPAPK MLGHELCRVC GDKASGFHYN VLSCEGCKGF FRRSVVRGGA
121 RRYACRGGGT CQMDAFMRRK CQQCRLRKCK EAGMREQCVL SEEQIRKKKI RKQQQESQSQ
181 SQSPVGPQGS SSSASGPGAS PGGSEAGSQG SGEGEGVQLT AAQELMIQQL VAAQLQCNKR
241 SFSDQPKVTP WPLGADPQSR DARQQRFAHF TELAIISVQE IVDFAKQVPG FLQLGREDQI
301 ALLKASTIEI MLLETARRYN HETECITFLK DFTYSKDDFH RAGLQVEFIN PIFEFSRAMR
361 RLGLDDAEYA LLIAINIFSA DRPNVQEPGR VEALQQPYVE ALLSYTRIKR PQDQLRFPRM
421 LMKLVSLRTL SSVHSEQVFA LRLQDKKLPP LLSEIWDVHE

Fig. 3 F:

1 atgtcctctc ctaccacgag ttccctggat accccctgc ctggaaatgg ccccctcag 61 cctggcgccc cttcttcttc acccactgta aaggaggagg gtccggagcc gtggcccggg 121 ggtccggacc ctgatgtccc aggcactgat gaggccagct cagcctgcag cacagactgg 181 gtcatcccag atcccgaaga ggaaccagag cgcaagcgaa agaagggccc agccccgaag 241 atgctgggcc acgagetttg ccgtgtctgt ggggacaagg cctccggett ccactacaac 301 gtgctcagct gcgaaggctg caagggcttc ttccggcgca gtgtggtccg tggtggggcc 361 aggcgctatg cctgccgggg tggcggaacc tgccagatgg acgctttcat gcggcgcaag 421 tgccagcagt gccggctgcg caagtgcaag gaggcaggga tgagggagca gtgcgtcctt 481 tctgaagaac agatccggaa gaagaagatt cggaaacagc agcaggagtc acagtcacag 541 tegeagteae etgtggggee geagggeage ageageteag cetetgggee tggggettee 601 cctggtggat ctgaggcagg cagccagggc tccggggaag gcgagggtgt ccagctaaca 661 gcggctcaag aactaatgat ccagcagttg gtggcggccc aactgcagtg caacaaacgc 721 teetteteeg accageceaa agteaegeee tggeecetgg gegeagaeee eeagteeega 781 gatgcccgcc agcaacgctt tgcccacttc acggagctgg ccatcatctc agtccaggag 841 atcgtggact tcgctaagca agtgcctggt ttcctgcagc tgggccggga ggaccagatc 901 gccctcctga aggcatccac tatcgagatc atgctgctag agacagccag gcgctacaac 961 cacgagacag agtgtatcac cttcttgaag gacttcacct acagcaagga cgacttccac 1021 cgtgcaggcc tgcaggtgga gttcatcaac cccatcttcg agttctcgcg ggccatgcgg 1081 cggctgggcc tggacgacgc tgagtacgcc ctgctcatcg ccatcaacat cttctcggcc 1141 gaccggccca acgtgcagga gccgggccgc gtggaggcgt tgcagcagcc ctacgtggag 1201 gcgctgctgt cctacacgcg catcaagagg ccgcaggacc agctgcgctt cccgcgcatg 1261 ctcatgaagc tggtgagcct gcgcacgctg agctctgtgc actcggagca ggtcttcgcc 1321 ttgcggctcc aggacaagaa gctgccgcct ctgctgtcgg agatctggga cgtccacgag 1381 tga

Fig. 4 A:

MOLNAME	MOLECULE STRUCTURE	EC50 AVG	EFFIC
		0.50	AVG
LN0000003252	N N CH ₃ CH ₃	0.59	110
LN0000007459	0 –	0.12	129
EN0000007439	CI CH ₃ CH ₃ CH ₃	0.12	129
LN0000011283	Br N CH ₃ CH ₃	0.18	137
TR1040007465	H H H	1.2	114
LN000007460		2,6	111
LN0000006500		0,34	133

	Chiral		
LN0000006494	Chiral	0,23	112

Fig. 4 A (cont.)

MOLNAME	MOLECULE STRUCTURE	EC50 AVG	EFFIC AVG
LN000007364		1.5	101
LN0000003492	N N N N N N N N N N N N N N N N N N N	0.11	115
LN0000007180		0.15	128
LN0000007179		0.78	127

Fig. 4 A (cont.):

Fig. 5:

Fig. 6

Liver X receptor alpha, LXRα (NM_005693) Cholesterol 7 α hydroxylase, Cyp7A1 (NM_000780)

FAS (NM_004104)

Stearyl CoA desaturase, SCD (XM 030447)

Sterol Response Element Binding Protein 1C, SREBP-1C (NM_004176)

ATP binding cassette transporter A1; ABCA1 (NM_005502)

ATP binding cassette transporter G1; ABCG1 (XM_032950)

ATP binding cassette transporter A1; ABCG5 (NM_031884)

ATP binding cassette transporter A1; ABCG8 AF324494

Apolipoprotein E, apoE (NM_000041)

Apolipoprotein C-I, apoC-I (NM_001645) Apolipoprotein C-II apoC-II (NM_000483)

Apoliporprotein C-IV, apoC-IV (U32576)

Lipoprotein Lipase, LPL (M15856) Cholesteryl Ester Transfer Protein, CETP (NM 000078)

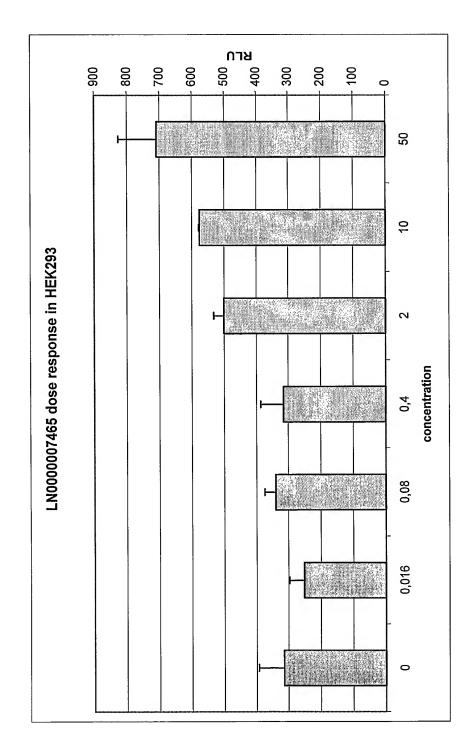


Fig. 7:

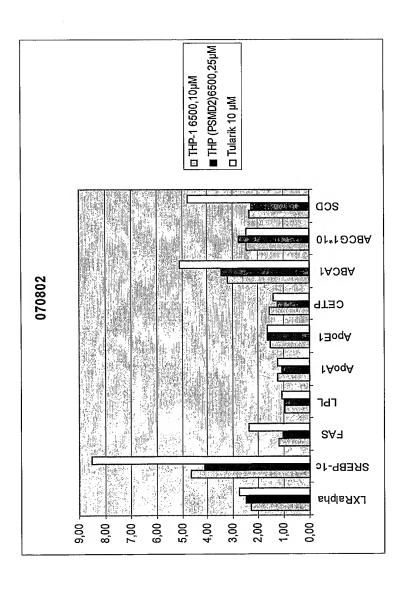


Fig. 8A:

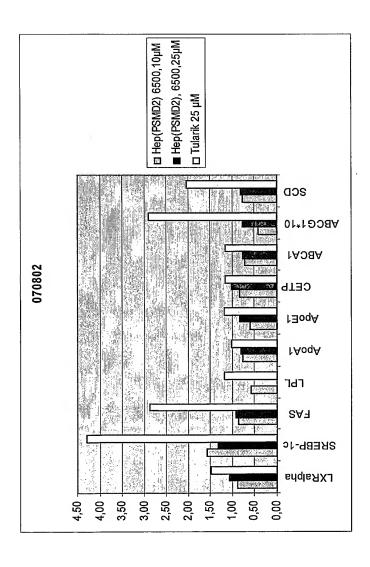


Fig. 8B

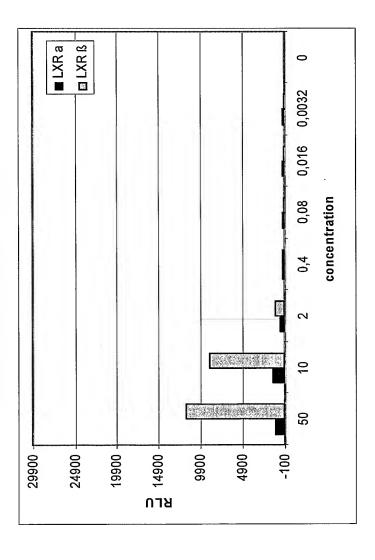
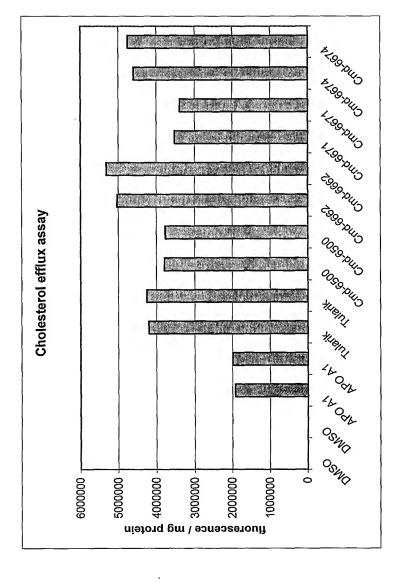


Fig. 9



19. TO:

1/15

SEQUENCE LISTING <110> LION bioscience <120> LXR Nuclear receptor binding compounds <130> L30281PCT <160> 8 <170> PatentIn version 3.1 <210> 1 <211> 52 <212> DNA <213> Homo sapiens <400> 1 ggggaccact ttgtacaaga aagctgggtc cccttctcag tctgttccac tt 52 <210> 2 <211> 1344 <212> DNA <213> Homo sapiens <400> 2 atgtccttqt qqctqqqqqc ccctqtqcct qacattcctc ctgactctqc qqtqqaqctq 60 tggaagccag gcgcacagga tgcaagcagc caggcccagg gaggcagcag ctgcatcctc 120 agagaggaag ccaggatgcc ccactctgct gggggtactg caggggtggg gctggaggct 180 gcagagccca cagccctgct caccagggca gagccccctt cagaacccac agagatccgt 240 300 ccacaaaagc ggaaaaaggg gccagcccc aaaatgctgg ggaacgagct atgcagcgtg 360 tgtggggaca aggcctcggg cttccactac aatgttctga gctgcgaggg ctgcaaggga ttcttccgcc gcagcgtcat caagggagcg cactacatct gccacagtgg cggccactgc 420 480 cccatggaca cctacatgcg tcgcaagtgc caggagtgtc ggcttcgcaa atgccgtcag gctggcatgc gggaggagtg tgtcctgtca gaagaacaga tccgcctgaa gaaactgaag 540 600 cggcaagagg aggaacaggc tcatgccaca tccttgcccc ccaggcgttc ctcaccccc caaatcctgc cccagctcag cccggaacaa ctgggcatga tcgagaagct cgtcgctgcc 660 720 cagcaacagt gtaaccggcg ctccttttct gaccggcttc gagtcacgcc ttggcccatg 780 gcaccagate eccatageeg ggaggeeegt cageageget ttgeeeactt caetgagetg gccatcgtct ctgtgcagga gatagttgac tttgctaaac agctacccgg cttcctgcag 840 ctcagccggg aggaccagat tgccctgctg aagacctctg cgatcgaggt gatgcttctg 900 gagacatete ggaggtacaa ecetgggagt gagagtatea eetteeteaa ggattteagt 960 tataaccggg aagactttgc caaagcaggg ctgcaagtgg aattcatcaa ccccatcttc 1020 gagttctcca gggccatgaa tgagctgcaa ctcaatgatg ccgagtttgc cttgctcatt 1080 gctatcagca tcttctctgc agaccggccc aacgtgcagg accagctcca ggtggagagg 1140

ctgcagcaca catatgtgga agccctgcat gcctacgtct ccatcacca tccccatgac

cgactgatgt tcccacggat gctaatgaaa ctggtgagcc tccggaccct gagcagcgtc 1260
cactcagagc aagtgtttgc actgcgtctg caggacaaaa agctcccacc gctgctctct 1320
gagatctggg atgtgcacga atga

<210> 3

<211> 1263

<212> PRT

<213> Homo sapiens

<400> 3

Met Leu Val Lys Pro Leu Pro Asp Ser Glu Glu Glu Gly His Asp Asn 1 $$ 5 $$ 10 $$ 15

Gln Glu Ala His Gln Lys Tyr Glu Thr Met Gln Cys Phe Ala Val Ser 20 25 30

Gln Pro Lys Ser Ile Lys Glu Glu Glu Glu Asp Leu Gln Ser Cys Leu 35 40 45

Ile Cys Val Ala Arg Arg Val Pro Met Lys Glu Arg Pro Val Leu Pro 50 55 60

Ser Ser Glu Ser Phe Thr Thr Arg Gln Asp Leu Gln Gly Lys Ile Thr 65 70 75 80

Ser Leu Asp Thr Ser Thr Met Arg Ala Ala Met Lys Pro Gly Trp Glu 85 90 95

Asp Leu Val Arg Arg Cys Ile Gln Lys Phe His Ala Gln His Glu Gly 100 105 110

Glu Ser Val Ser Tyr Ala Lys Arg His His His Glu Val Leu Arg Gln 115 120 125

Gly Leu Ala Phe Ser Gln Ile Tyr Arg Phe Ser Leu Ser Asp Gly Thr 130 135 140

Leu Val Ala Ala Gln Thr Lys Ser Lys Leu Ile Arg Ser Gln Thr Thr 145 150 155 160

Asn Glu Pro Gln Leu Val Ile Ser Leu His Met Leu His Arg Glu Gln
165 170 175

Asn Val Cys Val Met Asn Pro Asp Leu Thr Gly Gln Thr Met Gly Lys 180 185 190

Pro Leu Asn Pro Ile Ser Ser Asn Ser Pro Ala His Gln Ala Leu Cys 195 200 205

Ser	Gly 210	Asn	Pro	Gly	Gln	Asp 215	Met	Thr	Leu	Ser	Ser 220	Asn	Ile	Asn	Phe
Pro 225	Ile	Asn	Gly	Pro	Lys 230	Glu	Gln	Met	Gly	Met 235	Pro	Met	Gly	Arg	Phe 240
Gly	Gly	Ser	Gly	Gly 245	Met	Asn	His	Val	Ser 250	Gly	Met	Gln	Ala	Thr 255	Thr
Pro	Gln	Gly	Ser 260	Asn	Tyr	Ala	Leu	Lys 265	Met	Asn	Ser	Pro	Ser 270	Gln	Ser
Ser	Pro	Gly 275	Met	Asn	Pro	Gly	Gln 280	Pro	Thr	Ser	Met	Leu 285	Ser	Pro	Arg
His	Arg 290	Met	Ser	Pro	Gly	Val 295	Ala	Gly	Ser	Pro	Arg 300	Ile	Pro	Pro	Ser
Gln 305	Phe	Ser	Pro	Ala	Gly 310	Ser	Leu	His	Ser	Pro 315	Val	Gly	Val	Cys	Ser 320
Ser	Thr	Gly	Asn	Ser 325	His	Ser	Tyr	Thr	Asn 330	Ser	Ser	Leu	Asn	Ala 335	Leu
Gln	Ala	Leu	Ser 340	Glu	Gly	His	Gly	Val 345	Ser	Leu	Gly	Ser	Ser 350	Leu	Ala
Ser	Pro	Asp 355	Leu	Lys	Met	Gly	Asn 360	Leu	Gln	Asn	Ser	Pro 365	Val	Asn	Met
Asn	Pro 370	Pro	Pro	Leu	Ser	Lys 375	Met	Gly	Ser	Leu	Asp 380	Ser	Lys	Asp	Cys
Phe 385	Gly	Leu	Tyr	Gly	Glu 390	Pro	Ser	Glu	Gly	Thr 395	Thr	Gly	Gln	Ala	Glu 400
Ser	Ser	Суз	His	Pro 405	Gly	Glu	Gln	Lys	Glu 410	Thr	Asn	Asp	Pro	Asn 415	Leu
Pro	Pro	Ala	Val 420	Ser	Ser	Glu	Arg	Ala 425	Asp	Gly	Gln	Ser	Arg 430	Leu	His
Asp	Ser	Lys 435	Gly	Gln	Thr	Lys	Leu 440	Leu	Gln	Leu	Leu	Thr 445	Thr	Lys	Ser
Asp	Gln	Met	Glu	Pro	Ser	Pro	Leu	Ala	Ser	Ser	Leu	Ser	Asp	Thr	Asn

460

	4/

450

	450					455					400				
Lys 465	Asp	Ser	Thr	Gly	Ser 470	Leu	Pro	Gly	Ser	Gly 475	Ser	Thr	His	Gly	Thr 480
Ser	Leu	Lys	Glu	Lys 485	His	Lys	Ile	Leu	His 490	Arg	Leu	Leu	Gln	Asp 495	Ser
Ser	Ser	Pro	Val 500	Asp	Leu	Ala	Lys	Leu 505	Thr	Ala	Glu	Ala	Thr 510	Gly	Lys
Asp	Leu	Ser 515	Gln	Glu	Ser	Ser	Ser 520	Thr	Ala	Pro	Gly	Ser 525	Glu	Val	Thr
Ile	Lys 530	Gln	Glu	Pro	Val	Ser 535	Pro	Lys	Lys	Lys	Glu 540	Asn	Ala	Leu	Leu
Arg 545	Tyr	Leu	Leu	Asp	Lys 550	Asp	Asp	Thr	Lys	Asp 555	Ile	Gly	Leu	Pro	Glu 560
Ile	Thr	Pro	Lys	Leu 565	Glu	Arg	Leu	Asp	Ser 570	Lys	Thr	Asp	Pro	Ala 575	Ser
Asn	Thr	Lys	Leu 580	Ile	Ala	Met	Lys	Thr 585	Glu	Lys	Glu	Glu	Met 590	Ser	Phe
Glu	Pro	Gly 595	Asp	Gln	Pro	Gly	Ser 600	Glu	Leu	Asp	Asn	Leu 605	Glu	Glu	Ile
Leu	Asp 610	Asp	Leu	Gln	Asn	Ser 615	Gln	Leu	Pro	Gln	Leu 620	Phe	Pro	Asp	Thr
Arg 625	Pro	Gly	Ala	Pro	Ala 630	Gly	Ser	Val	Asp	Lys 635	Gln	Ala	Ile	Ile	Asn 640
Asp	Leu	Met	Gln	Leu 645	Thr	Ala	Glu	Asn	Ser 650	Pro	Val	Thr	Pro	Val 655	Gly
Ala	Gln	Lys	Thr 660	Ala	Leu	Arg	Ile	Ser 665	Gln	Ser	Thr	Phe	Asn 670	Asn	Pro
Arg	Pro	Gly 675	Gln	Leu	Gly	Arg	Leu 680	Leu	Pro	Asn	Gln	Asn 685	Leu	Pro	Leu
Asp	Ile 690	Thr	Leu	Gln	Ser	Pro 695	Thr	Gly	Ala	Gly	Pro 700	Phe	Pro	Pro	Ile

Arg Asn Ser Ser Pro Tyr Ser Val Ile Pro Gln Pro Gly Met Met Gly 705 710 715 Asn Gln Gly Met Ile Gly Asn Gln Gly Asn Leu Gly Asn Ser Ser Thr 730 Gly Met Ile Gly Asn Ser Ala Ser Arg Pro Thr Met Pro Ser Gly Glu 745 Trp Ala Pro Gln Ser Ser Ala Val Arg Val Thr Cys Ala Ala Thr Thr 760 Ser Ala Met Asn Arg Pro Val Gln Gly Gly Met Ile Arg Asn Pro Ala 770 Ala Ser Ile Pro Met Arg Pro Ser Ser Gln Pro Gly Gln Arg Gln Thr 785 790 795 800 Leu Gln Ser Gln Val Met Asn Ile Gly Pro Ser Glu Leu Glu Met Asn 805 810 Met Gly Gly Pro Gln Tyr Ser Gln Gln Gln Ala Pro Pro Asn Gln Thr 825 Ala Pro Trp Pro Glu Ser Ile Leu Pro Ile Asp Gln Ala Ser Phe Ala 835 840 Ser Gln Asn Arg Gln Pro Phe Gly Ser Ser Pro Asp Asp Leu Leu Cys 850 855 Pro His Pro Ala Ala Glu Ser Pro Ser Asp Glu Gly Ala Leu Leu Asp 865 870 Gln Leu Tyr Leu Ala Leu Arg Asn Phe Asp Gly Leu Glu Glu Ile Asp 885 Arg Ala Leu Gly Ile Pro Glu Leu Val Ser Gln Ser Gln Ala Val Asp 900 905 Pro Glu Gln Phe Ser Ser Gln Asp Ser Asn Ile Met Leu Glu Gln Lys 920 Ala Pro Val Phe Pro Gln Gln Tyr Ala Ser Gln Ala Gln Met Ala Gln 930 935 Gly Ser Tyr Ser Pro Met Gln Asp Pro Asn Phe His Thr Met Gly Gln

- Arg Pro Ser Tyr Ala Thr Leu Arg Met Gln Pro Arg Pro Gly Leu Arg 965 970 975
- Pro Thr Gly Leu Val Gln Asn Gln Pro Asn Gln Leu Arg Leu Gln Leu 980 985 990
- Gln His Arg Leu Gln Ala Gln Gln Asn Arg Gln Pro Leu Met Asn Gln 995 1000 1005
- Ile Ser Asn Val Ser Asn Val Asn Leu Thr Leu Arg Pro Gly Val 1010 1015 1020
- Pro Thr Gln Ala Pro Ile Asn Ala Gln Met Leu Ala Gln Arg Gln 1025 1030 1035
- Gln Gln Gln Val Gln Gln Arg Thr Leu Met Met Arg Gly Gln Gly 1055 1060 1065
- Leu Asn Met Thr Pro Ser Met Val Ala Pro Ser Gly Met Pro Ala 1070 $$ 1075 $$ 1080
- Thr Met Ser Asn Pro Arg Ile Pro Gln Ala Asn Ala Gln Gln Phe 1085 1090 1095
- Pro Phe Pro Pro Asn Tyr Gly Ile Ser Gln Gln Pro Asp Pro Gly 1100 $$ 1110
- Phe Thr Gly Ala Thr Thr Pro Gln Ser Pro Leu Met Ser Pro Arg 1115 1120 1125
- Met Ala His Thr Gln Ser Pro Met Met Gln Gln Ser Gln Ala Asn 1130 1135 1140
- Pro Ala Tyr Gln Ala Pro Ser Asp Ile Asn Gly Trp Ala Gln Gly 1145 1150 1155
- Asn Met Gly Gly Asn Ser Met Phe Ser Gln Gln Ser Pro Pro His 1160 1165 1170
- Phe Gly Gln Gln Ala Asn Thr Ser Met Tyr Ser Asn Asn Met Asn 1175 1180 1185
- Ile Asn Val Ser Met Ala Thr Asn Thr Gly Gly Met Ser Ser Met 1190 1200

Asn Gln Met Thr Gly Gln Ile Ser Met Thr Ser Val Thr Ser Val 1205 1210 1215

Pro Thr Ser Gly Leu Ser Ser Met Gly Pro Glu Gln Val Asn Asp 1220 1230

Pro Ala Leu Arg Gly Gly Asn Leu Phe Pro Asn Gln Leu Pro Gly 1235 1240 1245

Met Asp Met Ile Lys Gln Glu Gly Asp Thr Thr Arg Lys Tyr Cys 1250 1260

<210> 4 <211> 6158 <212> DNA <213> Homo sapiens

<213> Homo sapiens

<400> 4 ggcggccgca gcctcggcta cagcttcggc ggcqaaqqtc aqcqccqacq qcaqccqqca 60 cctgacggcg tgaccgaccc gagccgattt ctcttggatt tggctacaca cttatagatc 120 ttctgcactg tttacaggca cagttgctga tatgtgttca agatgagtgg gatgggagaa 180 aatacctctg acccctccag ggcagagaca agaaagcgca aggaatgtcc tgaccaactt 240 ggacccagcc ccaaaaggaa cactgaaaaa cgtaatcgtg aacaggaaaa taaatatata 300 gaagaacttg cagagttgat ttttgcaaat tttaatgata tagacaactt taacttcaaa 360 cctgacaaat gtgcaatctt aaaagaaact gtgaagcaaa ttcgtcagat caaagaacaa 420 gagaaagcag cagctgccaa catagatgaa gtgcagaagt cagatgtatc ctctacaggg 480 cagggtgtca tcgacaagga tgcgctgggg cctatgatqc ttqagqccct tgatqqqttc 540 ttctttgtag tgaacctgga aggcaacgtt gtgtttgtgt cagagaatgt gacacagtat 600 ctaaggtata accaagaaga gctgatgaac aaaagtgtat atagcatctt gcatgttggg 660 gaccacacgg aatttgtcaa aaacctgctg ccaaagtcta taggtaaatg ggggatcttq 720 gtctggcgaa cctccgaggc ggaacagcca taccttcaat tgtcggatgc tggtaaaacc 780 tttacctgat tcagaagagg agggtcatga taaccaggaa gctcatcaga aatatgaaac 840 tatgcagtgc ttcgctgtct ctcaaccaaa gtccatcaaa gaagaaggag aagatttgca 900 gtcctgcttg atttgcgtgg caagaagagt tcccatgaag gaaagaccag ttcttccctc 960 atcagaaagt tttactactc gccaggatct ccaaggcaag atcacgtctc tggataccag 1020 caccatgaga gcagccatga aaccaggctg ggaggacctg gtaagaaggt gtattcagaa 1080 gttccatgcg cagcatgaag gagaatctgt gtcctatgct aagaggcatc atcatgaagt 1140 actgagacaa ggattggcat tcagtcaaat ctatcgtttt tccttgtctg atggcactct 1200

tgttgctgca	caaacgaaga	gcaaactcat	ccgttctcag	actactaatg	aacctcaact	1260
tgtaatatct	ttacatatgc	ttcacagaga	gcagaatgtg	tgtgtgatga	atccggatct	1320
gactggacaa	acgatgggga	agccactgaa	tccaattagc	tctaacagcc	ctgcccatca	1380
ggccctgtgc	agtgggaacc	caggtcagga	catgaccctc	agtagcaata	taaattttcc	1440
cataaatggc	ccaaaggaac	aaatgggcat	gcccatgggc	aggtttggtg	gttctggggg	1500
aatgaaccat	gtgtcaggca	tgcaagcaac	cactcctcag	ggtagtaact	atgcactcaa	1560
aatgaacagc	ccctcacaaa	gcagccctgg	catgaatcca	ggacagccca	cctccatgct	1620
ttcaccaagg	catcgcatga	gccctggagt	ggctggcagc	cctcgaatcc	cacccagtca	1680
gttttcccct	gcaggaagct	tgcattcccc	tgtgggagtt	tgcagcagca	caggaaatag	1740
ccatagttat	accaacagct	ccctcaatgc	acttcaggcc	ctcagcgagg	ggcacggggt	1800
ctcattaggg	tcatcgttgg	cttcaccaga	cctaaaaatg	ggcaatttgc	aaaactcccc	1860
agttaatatg	aatcctcccc	cactcagcaa	gatgggaagc	ttggactcaa	aagactgttt	1920
tggactatat	ggggagccct	ctgaaggtac	aactggacaa	gcagagagca	gctgccatcc	1980
tggagagcaa	aaggaaacaa	atgaccccaa	cctgcccccg	gccgtgagca	gtgagagagc	2040
tgacgggcag	agcagactgc	atgacagcaa	agggcagacc	aaactcctgc	agctgctgac	2100
caccaaatct	gatcagatgg	agccctcgcc	cttagccagc	tctttgtcgg	atacaaacaa	2160
agactccaca	ggtagcttgc	ctggttctgg	gtctacacat	ggaacctcgc	tcaaggagaa	2220
gcataaaatt	ttgcacagac	tcttgcagga	cagcagttcc	cctgtggact	tggccaagtt	2280
aacagcagaa	gccacaggca	aagacctgag	ccaggagtcc	agcagcacag	ctcctggatc	2340
agaagtgact	attaaacaag	agccggtgag	ccccaagaag	aaagagaatg	cactacttcg	2400
ctatttgcta	gataaagatg	atactaaaga	tattggttta	ccagaaataa	ccccaaact	2460
tgagagactg	gacagtaaga	cagatcctgc	cagtaacaca	aaattaatag	caatgaaaac	2520
tgagaaggag	gagatgagct	ttgagcctgg	tgaccagcct	ggcagtgagc	tggacaactt	2580
ggaggagatt	ttggatgatt	tgcagaatag	tcaattacca	cagcttttcc	cagacacgag	2640
gccaggcgcc	cctgctggat	cagttgacaa	gcaagccatc	atcaatgacc	tcatgcaact	2700
cacagctgaa	aacagccctg	tcacacctgt	tggagcccag	aaaacagcac	tgcgaatttc	2760
acagagcact	tttaataacc	cacgaccagg	gcaactgggc	aggttattgc	caaaccagaa	2820
tttaccactt	gacatcacat	tgcaaagccc	aactggtgct	ggacctttcc	caccaatcag	2880
aaacagtagt	ccctactcag	tgatacctca	gccaggaatg	atgggtaatc	aagggatgat	2940
aggaaaccaa	ggaaatttag	ggaacagtag	cacaggaatg	attggtaaca	gtgcttctcg	3000
gcctactatg	ccatctggag	aatgggcacc	gcagagttcg	gctgtgagag	tcacctgtgc	3060

tgctaccacc	agtgccatga	accggccagt	ccaaggaggt	atgattcgga	acccagcagc	3120
cagcatcccc	atgaggccca	gcagccagcc	tggccaaaga	cagacgcttc	agtctcaggt	3180
catgaatata	gggccatctg	aattagagat	gaacatgggg	ggacctcagt	atagccaaca	3240
acaagctcct	ccaaatcaga	ctgccccatg	gcctgaaagc	atcctgccta	tagaccaggc	3300
gtcttttgcc	agccaaaaca	ggcagccatt	tggcagttct	ccagatgact	tgctatgtcc	3360
acatcctgca	gctgagtctc	cgagtgatga	gggagctctc	ctggaccagc	tgtatctggc	3420
cttgcggaat	tttgatggcc	tggaggagat	tgatagagcc	ttaggaatac	ccgaactggt	3480
cagccagagc	caagcagtag	atccagaaca	gttctcaagt	caggattcca	acatcatgct	3540
ggagcagaag	gcgcccgttt	tcccacagca	gtatgcatct	caggcacaaa	tggcccaggg	3600
tagctattct	cccatgcaag	atccaaactt	tcacaccatg	ggacagcggc	ctagttatgc	3660
cacactccgt	atgcagccca	gaccgggcct	caggcccacg	ggcctagtgc	agaaccagcc	3720
aaatcaacta	agacttcaac	ttcagcatcg	cctccaagca	cagcagaatc	gccagccact	3780
tatgaatcaa	atcagcaatg	tttccaatgt	gaacttgact	ctgaggcctg	gagtaccaac	3840
acaggcacct	attaatgcac	agatgctggc	ccagagacag	agggaaatcc	tgaaccagca	3900
tcttcgacag	agacaaatgc	atcagcaaca	gcaagttcag	caacgaactt	tgatgatgag	3960
aggacaaggg	ttgaatatga	caccaagcat	ggtggctcct	agtggtatgc	cagcaactat	4020
gagcaaccct	cggattcccc	aggcaaatgc	acagcagttt	ccatttcctc	caaactacgg	4080
aataagtcag	caacctgatc	caggctttac	tggggctacg	actccccaga	gcccacttat	4140
gtcaccccga	atggcacata	cacagagtcc	catgatgcaa	cagtctcagg	ccaacccagc	4200
ctatcaggcc	ccctccgaca	taaatggatg	ggcgcagggg	aacatgggcg	gaaacagcat	4260
gttttcccag	cagtccccac	cacactttgg	gcagcaagca	aacaccagca	tgtacagtaa	4320
caacatgaac	atcaatgtgt	ccatggcgac	caacacaggt	ggcatgagca	gcatgaacca	4380
gatgacagga	cagatcagca	tgacctcagt	gacctccgtg	cctacgtcag	ggctgtcctc	4440
catgggtccc	gagcaggtta	atgatcctgc	tctgagggga	ggcaacctgt	tcccaaacca	4500
gctgcctgga	atggatatga	ttaagcagga	gggagacaca	acacggaaat	attgctgaca	4560
ctgctgaagc	cagttgcttc	ttcagctgac	cgggctcact	tgctcaaaac	acttccagtc	4620
tggagagctg	tgtctatttg	tttcaaccca	actgacctgc	cagccggttc	tgctagagca	4680
gacaggcctg	gccctggttc	ccagggtggc	gtccactcgg	ctgtggcagg	aggagctgcc	4740
tcttctcttg	acagtctgaa	gctcgcatcc	agacagtcgc	tcagtctgtt	cactgcattc	4800
accttagtgc	aacttagatc	tctcctgcaa	aagtaaatgt	tgacaggcaa	atttcatacc	4860
catgtcagat	tgaatgtatt	taaatgtatg	tatttaagga	gaaccatgct	cttgttctgt	4920
tcctgttcgg	ttccagacac	tggtttcttg	ctttgttttc	cctggctaac	agtctagtgc	4980

aaaagattaa	gattttatct	gggggaaaga	aaagaatttt	ttaaaaaatt	aaactaaaga	5040
tgttttaagc	taaagcctga	atttgggatg	gaagcaggac	agacaccgtg	gacagcgctg	5100
tatttacaga	cacacccagt	gcgtgaagac	caacaaagtc	acagtcgtat	ctctagaaag	5160
ctctaaagac	catgttggaa	agagtctcca	gttactgaac	agatgaaaag	gagcctgtga	5220
gagggctgtt	aacattagca	aatattttt	ccttgttttt	tctttgttaa	aaccaaactg	5280
gttcacctga	atcatgaatt	gagaagaaat	aattttcatt	tctaaattaa	gtccctttta	5340
gtttgatcag	acagcttgaa	tcagcatctc	ttcttccctg	tcagcctgac	tcttcccttc	5400
ccctctctca	ttccccatac	tccctatttt	cattcctttt	ttaaaaaata	atataagcta	5460
cagaaaccag	gtaagccctt	tatttcctta	aatgttttgc	cagccactta	ccaattgcta	5520
agtattgaat	ttcagaaaaa	aaaaatgcat	ttactggcaa	ggagaagagc	aaagttaagg	5580
cttgatacca	atcgagctaa	ggatacctgc	tttggaagca	tgtttattct	gttccccagc	5640
aactctggcc	tccaaaatgg	gagaaaacgc	cagtgtgttt	aaattgatag	cagatatcac	5700
gacagattta	acctctgcca	tgtgttttt	attttgtttt	ttagcagtgc	tgactaagcc	5760
gaagttttgt	aaggtacata	aaatccaatt	tatatgtaaa	caagcaataa	tttaagttga	5820
gaacttatgt	gttttaattg	tataattttt	gtgaggtata	catattgtgg	aattgactca	5880
aaaatgaggt	acttcagtat	taaattagat	atcttcatag	caatgtctcc	taaaggtgtt	5940
ttgtaaagga	tatcaatgcc	ttgattagac	ctaatttgta	gacttaagac	ttttattt	6000
ctaaaccttg	tgattctgct	tataagtcat	ttatctaatc	tatatgatat	gcagccgctg	6060
taggaaccaa	ttcttgattt	ttatatgttt	atattctttc	ttaatgaacc	ttagaaagac	6120
tacatgttac	taagcaggcc	acttttatgg	ttgttttt			6158

<210> 5 <211> 460 <212> PRT

<213> Homo sapiens <400> 5

Met Ser Ser Pro Thr Thr Ser Ser Leu Asp Thr Pro Leu Pro Gly Asn 1 5 10

Gly Pro Pro Gln Pro Gly Ala Pro Ser Ser Pro Thr Val Lys Glu

Glu Gly Pro Glu Pro Trp Pro Gly Gly Pro Asp Pro Asp Val Pro Gly 40

Thr Asp Glu Ala Ser Ser Ala Cys Ser Thr Asp Trp Val Ile Pro Asp

11/15

Pro Glu Glu Glu Pro Glu Arg Lys Arg Lys Lys Gly Pro Ala Pro Lys Met Leu Gly His Glu Leu Cys Arg Val Cys Gly Asp Lys Ala Ser Gly Phe His Tyr Asn Val Leu Ser Cys Glu Gly Cys Lys Gly Phe Phe Arg 105 Arg Ser Val Val Arg Gly Gly Ala Arg Arg Tyr Ala Cys Arg Gly Gly 115 120 Gly Thr Cys Gln Met Asp Ala Phe Met Arg Arg Lys Cys Gln Gln Cys 135 Arg Leu Arg Lys Cys Lys Glu Ala Gly Met Arg Glu Gln Cys Val Leu Ser Glu Glu Gln Ile Arg Lys Lys Lys Ile Arg Lys Gln Gln Glu 165 170 Ser Gln Ser Gln Ser Pro Val Gly Pro Gln Gly Ser Ser Ser 180 185 190 Ser Ala Ser Gly Pro Gly Ala Ser Pro Gly Gly Ser Glu Ala Gly Ser 195 200 205 Gln Gly Ser Gly Glu Gly Glu Gly Val Gln Leu Thr Ala Ala Gln Glu 210 215 220 Leu Met Ile Gln Gln Leu Val Ala Ala Gln Leu Gln Cys Asn Lys Arg 225 230 Ser Phe Ser Asp Gln Pro Lys Val Thr Pro Trp Pro Leu Gly Ala Asp 245 250

Pro Gln Ser Arg Asp Ala Arg Gln Gln Arg Phe Ala His Phe Thr Glu 265

Leu Ala Ile Ile Ser Val Gln Glu Ile Val Asp Phe Ala Lys Gln Val 275 280

Pro Gly Phe Leu Gln Leu Gly Arg Glu Asp Gln Ile Ala Leu Leu Lys 290 295

Ala Ser Thr Ile Glu Ile Met Leu Leu Glu Thr Ala Arg Arg Tyr Asn 305 310 315

His Glu Thr Glu Cys Ile Thr Phe Leu Lys Asp Phe Thr Tyr Ser Lys 325 330 335	
Asp Asp Phe His Arg Ala Gly Leu Gln Val Glu Phe Ile Asn Pro Ile 340 345 350	
Phe Glu Phe Ser Arg Ala Met Arg Arg Leu Gly Leu Asp Asp Ala Glu 355 360 365	
Tyr Ala Leu Leu Ile Ala Ile Asn Ile Phe Ser Ala Asp Arg Pro Asn 370 380	
Val Gln Glu Pro Gly Arg Val Glu Ala Leu Gln Gln Pro Tyr Val Glu 385 390 395 400	
Ala Leu Leu Ser Tyr Thr Arg Ile Lys Arg Pro Gln Asp Gln Leu Arg 405 410 415	
Phe Pro Arg Met Leu Met Lys Leu Val Ser Leu Arg Thr Leu Ser Ser 420 425 430	
Val His Ser Glu Gln Val Phe Ala Leu Arg Leu Gln Asp Lys Lys Leu 435 440 445	
Pro Pro Leu Leu Ser Glu Ile Trp Asp Val His Glu 450 455 460	
<210> 6 <211> 1383 <212> DNA <213> Homo sapiens	
<400> 6 atgtcctctc ctaccacgag ttccctggat acccccctgc ctggaaatgg cccccctcag 60	
cctggcgccc cttcttcttc acccactgta aaggaggagg gtccggagcc gtggcccggg 120	
ggtccggacc ctgatgtccc aggcactgat gaggccagct cagcctgcag cacagactgg 180	
gtcatcccag atcccgaaga ggaaccagag cgcaagcgaa agaagggccc agccccgaag 240	
atgctgggcc acgagetttg ccgtgtctgt ggggacaagg cctccggctt ccactacaac 300	
gtgctcagct gcgaaggctg caagggcttc ttccggcgca gtgtggtccg tggtggggcc 360	

aggcgctatg cctgccgggg tggcggaacc tgccagatgg acgctttcat gcggcgcaag

tgccagcagt gccggctgcg caagtgcaag gaggcaggga tgagggagca gtgcgtcctt

tctgaagaac agatccggaa gaagaagatt cggaaacagc agcaggagtc acagtcacag

tcgcagtcac ctgtggggcc gcagggcagc agcagctcag cctctgggcc tggggcttcc

420

480

540

cctggtggat	ctgaggcagg	cagccagggc	tccggggaag	gcgagggtgt	ccagctaaca	660
gcggctcaag	aactaatgat	ccagcagttg	gtggcggccc	aactgcagtg	caacaaacgc	720
teetteteeg	accagcccaa	agtcacgccc	tggcccctgg	gcgcagaccc	ccagtcccga	780
gatgcccgcc	agcaacgctt	tgcccacttc	acggagctgg	ccatcatctc	agtccaggag	840
atcgtggact	tcgctaagca	agtgcctggt	ttcctgcagc	tgggccggga	ggaccagatc	900
gccctcctga	aggcatccac	tatcgagatc	atgctgctag	agacagccag	gcgctacaac	960
cacgagacag	agtgtatcac	cttcttgaag	gacttcacct	acagcaagga	cgacttccac	1020
cgtgcaggcc	tgcaggtgga	gttcatcaac	cccatcttcg	agttctcgcg	ggccatgcgg	1080
cggctgggcc	tggacgacgc	tgagtacgcc	ctgctcatcg	ccatcaacat	cttctcggcc	1140
gaccggccca	acgtgcagga	gccgggccgc	gtggaggcgt	tgcagcagcc	ctacgtggag	1200
gcgctgctgt	cctacacgcg	catcaagagg	ccgcaggacc	agctgcgctt	cccgcgcatg	1260
ctcatgaagc	tggtgagcct	gcgcacgctg	agctctgtgc	actcggagca	ggtcttcgcc	1320
ttgcggctcc	aggacaagaa	gctgccgcct	ctgctgtcgg	agatctggga	cgtccacgag	1380
tga						1383

<210> 7 <211> 49 <212> DNA

<213> Homo sapiens

<400> 7

ggggacaagt ttgtacaaaa aagcaggctc gcttcgcaaa tgccgtcag

49

<210> 8 <211> 447 <212> PRT <213> Homo sapiens <400> 8

Met Ser Leu Trp Leu Gly Ala Pro Val Pro Asp Ile Pro Pro Asp Ser 1 5

Ala Val Glu Leu Trp Lys Pro Gly Ala Gln Asp Ala Ser Ser Gln Ala

Gln Gly Gly Ser Ser Cys Ile Leu Arg Glu Glu Ala Arg Met Pro His 40

Ser Ala Gly Gly Thr Ala Gly Val Gly Leu Glu Ala Ala Glu Pro Thr 55

Ala Leu Leu Thr Arg Ala Glu Pro Pro Ser Glu Pro Thr Glu Ile Arg 70 75

Pro Gln Lys Arg Lys Lys Gly Pro Ala Pro Lys Met Leu Gly Asn Glu

Leu Cys Ser Val Cys Gly Asp Lys Ala Ser Gly Phe His Tyr Asn Val 100 105 110

Leu Ser Cys Glu Gly Cys Lys Gly Phe Phe Arg Arg Ser Val Ile Lys 115 120 125

Gly Ala His Tyr Ile Cys His Ser Gly Gly His Cys Pro Met Asp Thr 130 135 140

Tyr Met Arg Arg Lys Cys Gln Glu Cys Arg Leu Arg Lys Cys Arg Gln 145 150 155 160

Ala Gly Met Arg Glu Glu Cys Val Leu Ser Glu Glu Gln Ile Arg Leu 165 170 175

Lys Lys Leu Lys Arg Gln Glu Glu Glu Gln Ala His Ala Thr Ser Leu 180 185 190

Pro Pro Arg Arg Ser Ser Pro Pro Gln Ile Leu Pro Gln Leu Ser Pro 195 200 205

Glu Gln Leu Gly Met Ile Glu Lys Leu Val Ala Ala Gln Gln Gln Cys 210 215 220

Asn Arg Arg Ser Phe Ser Asp Arg Leu Arg Val Thr Pro Trp Pro Met 225 230 235 . 240

Ala Pro Asp Pro His Ser Arg Glu Ala Arg Gln Gln Arg Phe Ala His 245 250 255

Phe Thr Glu Leu Ala Ile Val Ser Val Gln Glu Ile Val Asp Phe Ala 260 265 270

Lys Gln Leu Pro Gly Phe Leu Gln Leu Ser Arg Glu Asp Gln Ile Ala 275 280 285

Leu Leu Lys Thr Ser Ala Ile Glu Val Met Leu Leu Glu Thr Ser Arg 290 295 300

Arg Tyr Asn Pro Gly Ser Glu Ser Ile Thr Phe Leu Lys Asp Phe Ser 305 310 315 320

Tyr Asn Arg Glu Asp Phe Ala Lys Ala Gly Leu Gln Val Glu Phe Ile 325 330 335

Asn Pro Ile Phe Glu Phe Ser Arg Ala Met Asn Glu Leu Gln Leu Asn 340 345 350

- Asp Ala Glu Phe Ala Leu Leu Ile Ala Ile Ser Ile Phe Ser Ala Asp 355 360 365
- Arg Pro Asn Val Gln Asp Gln Leu Gln Val Glu Arg Leu Gln His Thr 370 375 380
- Tyr Val Glu Ala Leu His Ala Tyr Val Ser Ile His His Pro His Asp 385 390 390 395
- Arg Leu Met Phe Pro Arg Met Leu Met Lys Leu Val Ser Leu Arg Thr 405 410 415
- Leu Ser Ser Val His Ser Glu Gln Val Phe Ala Leu Arg Leu Gln Asp 420 425 430
- Lys Lys Leu Pro Pro Leu Leu Ser Glu Ile Trp Asp Val His Glu 435 440 445

Internat Application No PCT/EP 03/07067

CLASSIFICATION OF SUBJECT MATTER C 7 A61K31/517 C07D C07D239/95 CO7D401/12 C07D403/04 A61P3/06 According to International Patent Classification (IPC) or to both national classification and IPC **B. FIELDS SEARCHED** Minimum documentation searched (classification system followed by classification symbols) IPC 7 C07D A61K A61P Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched Electronic data base consulted during the international search (name of data base and, where practical, search terms used) EPO-Internal, WPI Data, CHEM ABS Data C. DOCUMENTS CONSIDERED TO BE RELEVANT Category 6 Citation of document, with indication, where appropriate, of the relevant passages Relevant to claim No. χ WO 02 062798 A (REDDY RESEARCH FOUNDATION) 1,13-3315 August 2002 (2002-08-15) page 106, line 22 - page 107, line 6 page 1, line 17 - page 2, line 16 χ WO 97 20823 A (CRISCIONE LEOLUCA 1,13,18, ;YAMAGUCHI YASUCHIKA (CH); CIBA GEIGY AG 22,30, (CH); M) 12 June 1997 (1997-06-12) 32,35 page 74; example 38 page 1, paragraph 1 χ WO 02 48152 A (BAKTHAVATCHALAM RAJAGOPAL 1,13,18, ;BRIELMANN HARRY L (US); ELLIOTT RICHARD) 22,30, 20 June 2002 (2002-06-20) 32,35 page 59; example 12 page 6, paragraph 17 -/-χ Further documents are listed in the continuation of box C. Patent family members are listed in annex. Special categories of cited documents: "T" later document published after the International filing date or priority date and not in conflict with the application but "A" document defining the general state of the art which is not considered to be of particular relevance cited to understand the principle or theory underlying the invention "E" earlier document but published on or after the international *X* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to filing date *L* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another involve an inventive step when the document is taken alone "Y" document of particular relevance; the claimed invention citation or other special reason (as specified) cannot be considered to involve an inventive step when the document is combined with one or more other such docudocument referring to an oral disclosure, use, exhibition or ments, such combination being obvious to a person skilled in the art. other means document published prior to the international filing date but later than the priority date claimed "&" document member of the same patent family Date of the actual completion of the international search Date of mailing of the international search report 22 September 2003 06/10/2003 Name and mailing address of the ISA Authorized officer European Patent Office, P.B. 5818 Patentlaan 2 NL – 2280 HV Rijswijk Tel. (+31–70) 340–2040, Tx. 31 651 epo nl, Fax: (+31–70) 340–3016 Kollmannsberger, M.

		PCI/EP 03/0/06/
	ation) DOCUMENTS CONSIDERED TO BE RELEVANT	
Category °	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
А	COLLINS, J. L. ET AL.: "Identification of a Nonsteroidal Liver X Receptor Agonist through Parallel Array Synthesis of Tertiary Amines" JOURNAL OF MEDICINAL CHEMISTRY, vol. 45, 2002, pages 1963-1966, XP002225147 cited in the application the whole document	1-33
X	GUPTA C M ET AL: "Drugs acting on the central nervous system. Syntheses of substituted quinazolones and quinazolines and triazepino- and triazocinoquinazolones" JOURNAL OF MEDICINAL CHEMISTRY, AMERICAN CHEMICAL SOCIETY. WASHINGTON, US, vol. 11, no. 2, 26 February 1968 (1968-02-26), pages 392-395, XP002156695 ISSN: 0022-2623 examples 13-16,22,23,38; table 2	1,4,8,13
X	MANABU HORI ET AL: "Novel 4-Substituted 2-Piperazinylquinazolines as potent Anticonvulsive and Antihypoxic Agents" CHEMICAL AND PHARMACEUTICAL BULLETIN, PHARMACEUTICAL SOCIETY OF JAPAN. TOKYO, JP, vol. 38, no. 5, 1990, pages 1286-1291, XP002128282 ISSN: 0009-2363 examples 3A-3H; table II	1
X	US 3 609 152 A (HESS HANS-JURGEN E ET AL) 28 September 1971 (1971-09-28) examples III-X	1,4,8,13
X	DATABASE CHEMCATS CHEMICAL ABSTRACTS SERVICE, COLUMBUS, OHIO, US; 2001, XP002225148 Order Number: TRG10400#07364-D; TRG10400#01891-D; TRG10400#01815-D; TRG10400#01814-D; TRG10400#01812-D; TRG10400#01811-D; TRG10400#01809-D; TRG10400#01736-D; TRG10400#01735-D; TRG10400#01732-D; TRG10400#01729-D & "Chem.Folio" 15 January 2001 (2001-01-15), LION BIOSCIENCE AG , HEIDELBERG, GERMANY	1-3,9

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)	
This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:	
1. X Claims Nos.: because they relate to subject matter not required to be searched by this Authority, namely:	
Claims 14-25, 34, 35 are directed to a method of treatment of the human/animal body. Insofar as the claims could be searched, the search has been carried out based on the alleged effects of the compounds.	
Claims Nos.: Claims Nos.: Claims Nos.: Claims Nos.: because they relate to parts of the International Application that do not comply with the prescribed requirements to such an extent that no meaningful International Search can be carried out, specifically:	
see FURTHER INFORMATION sheet PCT/ISA/210	
Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).	
Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)	\dashv
This International Searching Authority found multiple Inventions in this international application, as follows:	
1. As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.	
2. As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.	
3. As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos.:	
4. No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:	
Remark on Protest The additional search fees were accompanied by the applicant's protest. No protest accompanied the payment of additional search fees.	

FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

Continuation of Box I.2

Claims Nos.: -

Claims 1-12 encompass a large number of known compounds. The initial phase of the search revealed a very large number of documents relevant to the issue of novelty. So many documents were retrieved that it is impossible to determine which parts of the claim(s) may be said to define subject-matter for which protection might legitimately be sought (Article 6 PCT). Additionally, support within the meaning of Article 6 PCT and disclosure within the meaning of Article 5 PCT is to be found, for only a very small proportion of the compounds and methods claimed. For these reasons, a meaningful search over the whole breadth of the claims is impossible. Consequently, the search is only complete for:

Compounds according to claims 1-10 which are mentioned in the prior art to have useful properties in the treatment of the diseases mentioned in claims 29-32; compounds as such according to claims 4-10.

Only some documents relevant to other subject-matter of the claims have been cited for illustration.

The applicant's attention is drawn to the fact that claims relating to inventions in respect of which no international search report has been established need not be the subject of an international preliminary examination (Rule 66.1(e) PCT). The applicant is advised that the EPO policy when acting as an International Preliminary Examining Authority is normally not to carry out a preliminary examination on matter which has not been searched. This is the case irrespective of whether or not the claims are amended following receipt of the search report or during any Chapter II procedure. If the application proceeds into the regional phase before the EPO, the applicant is reminded that a search may be carried out during examination before the EPO (see EPO Guideline C-VI, 8.5), should the problems which led to the Article 17(2) declaration be overcome.



Patent document cited in search report	Publication date	Patent fa membe		Publication date
WO 02062798	A 15-08-2002	WO 0206	2798 A2 2799 A1 9175 A1	15-08-2002 15-08-2002 14-11-2002
WO 9720823	A 12-06-1997	WO 972	2996 A 0823 A2 0020 A	27-06-1997 12-06-1997 01-06-1997
WO 0248152	A 20-06-2002	WO 024	7602 A 8152 A2 6652 A1	24-06-2002 20-06-2002 20-02-2003
US 3609152	A 28-09-1971	DE 1629 FR 9 GB 1066 GB 1176	 8216 A 0127 A1 5267 M 2357 A 4272 A 4273 A	22-09-1966 12-03-1970 31-07-1967 22-03-1967 17-12-1969 17-12-1969